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LUIZ DANIEL DE BARROS

**AVALIAÇÃO DOS POMBOS DA ESPÉCIE *ZENaida*  
*AURICULATA* COMO HOSPEDEIRO INTERMEDIÁRIO DO  
*NEOSPORA CANINUM***

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Tese de doutorado apresentada ao Programa de Pós-graduação em Ciência Animal (área de concentração: Sanidade Animal) do Centro de Ciências Agrárias da Universidade Estadual de Londrina, como requisito para a obtenção do título de Doutor em Ciência Animal

Orientador: Prof. Dr. João Luis Garcia

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**BANCA EXAMINADORA**

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Londrina, 01 de abril de 2016.

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**You can't climb the ladder of success with  
your hands in your pockets**  
(Arnold Schwarzenegger)

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## RESUMO

*Neospora caninum* é um protozoário, parasita intracelular obrigatório pertencente filo Apicomplexa e família Sarcocystidae. O parasita possui distribuição mundial e é responsável pela neosporose, uma doença primária de cães e bovinos, sendo considerada como uma das principais causas de abortamento em bovinos. Cães e outros canídeos são os hospedeiros definitivos e eliminam oocistos no ambiente. Vários animais domésticos e silvestres, incluindo aves, tem sido descritos como hospedeiros intermediários, entretanto o papel desses animais no ciclo epidemiológico ainda é incerto. O objetivo do presente estudo foi avaliar se pombos da espécie *Zenaida auriculata* podem atuar como hospedeiros intermediários do *N. caninum*, contribuindo assim com a transmissão da doença. No primeiro experimento, 249 pombos *Z. auriculata* foram capturados em três diferentes áreas da cidade de Londrina, Paraná. Amostras de soro e cérebro foram coletadas para a realização dos testes sorológicos e molecular respectivamente. Por meio da técnica de ELISA, 31,72% (79/249) das amostras foram positivas, com maior número de aves positivas capturadas em uma indústria de processamento de grãos. Não foi observado DNA do parasita utilizando a técnica de PCR em nenhuma das amostras de tecido cerebral. No segundo experimento, 18 pombos foram divididos em 4 grupos, G1, G2 e G3, composto por 5 animais cada, receberam  $2 \times 10^6$  taquizoítos da cepa NC-1 do *N. caninum* pelas vias subcutânea, intramuscular e intraperitoneal respectivamente. G4, composto por 3 animais, permaneceu como grupo controle. A técnica de ELISA foi utilizada para verificar a cinética dos anticorpos contra o parasita após a inoculação. No sétimo e 14º dia, um animal dos grupos G1, G2 e G3 foram eutanasiados para detecção do parasita pela PCR. Após 28 dias, todos os outros animais foram eutanasiados e tecidos (cérebro, coração, fígado, pulmão, rim e músculo peitoral) foram coletados para análise de PCR e exame histopatológico. Foi realizado bioensaio em cães para verificar cistos na musculatura de pombos experimentalmente infectados. Quatro cães foram alimentados com tecidos de pombos inoculados previamente de somente um grupo. Exames coproparasitológico e imunológico foram realizados para avaliar a infecção nos cães. Todas as aves inoculadas soroconverteram após desafio, com maior produção de anticorpos nas aves inoculadas pela via intraperitoneal. Um animal do grupo 3 eutanaziado no 7 dia após inoculação foi positivo na PCR dos tecidos. Apenas um animal do G3 apresentou um infiltrado inflamatório mononuclear no pulmão e rim, compatível com neosporose no exame histopatológico. Nenhum cão soroconverteu ou eliminou oocistos durante o período experimental. Esse estudo verificou pela primeira vez a presença de anticorpos contra *N. caninum* em pombos *Z. auriculata* de vida livre, demonstrando que esses animais tiveram um contato prévio com o *N. caninum*. Nossos resultados também indicaram que a via intraperitoneal induziu infecção nos pombos, entretanto essas aves foram consideradas resistentes à infecção crônica. Estudos adicionais devem ser conduzidos, especialmente utilizando outros modelos de infecção, para avaliar a importância dos pombos *Z. auriculata* na epidemiologia da neosporose.

**Palavras-chave:** Neosporose. Pombo. Sorologia. ELISA. Bioensaio em cães.

BARROS, Luiz Daniel. **Evaluation of doves from *Zenaida auriculata* species as intermediate host of *Neospora caninum***. 2016. 80p. Thesis (Animal Science) – Universidade Estadual de Londrina, Londrina, 2016.

## ABSTRACT

*Neospora caninum* is a protozoan, obligate intracellular parasite that belongs to Apicomplexa phylum and Sarcocystidae family. The parasite has worldwide distribution and is responsible for neosporosis, a primarily in dogs and cattle, being considered as one of the major causes of abortion in cattle. Dogs and other canids are the definitive hosts and shed oocysts in the environment. Many domestic and wild animals, including birds, have been described as intermediate host, however, the role of these animals in epidemiological cycle is uncertain. The aim of the present study was to evaluate if doves from *Zenaida auriculata* species can act as intermediate host of *N. caninum*, contributing with transmission of the disease. In the first experiment, 249 doves *Z. auriculata* were captured in three different areas from Londrina city, Paraná. Serum and brain samples were collected to perform serological and molecular tests respectively. By means of ELISA, 31.72% (79/249) of samples were positive, with higher number of positive birds captured in an soybean seed plant. It was not observed DNA from the parasite using the PCR technique in all brain tissue samples tested. In the second experiment, 18 doves were divided in 4 groups, G1, G2 and G3, composed by 5 animals each, received  $2 \times 10^6$  tachyzoites from NC-1 strain of *N. caninum* by subcutaneous, intramuscular and intraperitoneal route respectively. G4, composed by 3 animals, remained as control group. ELISA technique was used to verify the kinetics of antibodies against the parasite after inoculation. On 7<sup>th</sup> and 14<sup>th</sup> day, one animal from G1, G2 and G3 groups, were euthanized to detect the parasite by PCR. After 28 days, all other animals were euthanized and tissue (brain, heart, liver, lung, kidney and pectoral muscle) were collected for PCR analysis and histopathological examination. Bioassay in dogs were performed to verify cysts in the musculature of doves infected experimentally. Four dogs were fed with doves tissue from only one group. Coproparasitological and immunological examination were performed to evaluate dogs infection. All birds inoculated seroconverted after challenge, with higher antibodies production in birds inoculated through intraperitoneal route. One animal from group 3 that was euthanized on 7<sup>th</sup> day after inoculation was positive in PCR from tissues. Only one animal from G3 showed a mononuclear inflammatory infiltrate in lung and kidney, which is compatible with neosporosis in the histopathology evaluation. None dogs seroconverted or shed oocysts during experimental period. This study verified for the first time the presence of antibodies against *N. caninum* in free-ranging doves *Z. auriculata*, showing that these animals had previously contact with *N. caninum*. Our results also indicated that intraperitoneal route induced infection in doves, however these birds were considered resistant to chronic infection. Additional studies should be conducted, especially using other infection models, to evaluate the importance of *Z. auriculata* doves in the epidemiology of neosporosis.

**Keywords:** Neosporosis. Eared doves. Serology. ELISA. Dogs bioassay.

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## 1 INTRODUÇÃO

A neosporose é uma doença infecciosa de origem parasitária causada pelo protozoário *Neospora caninum* (DUBEY et al., 1988). A doença é de grande importância na bovinocultura mundial, devido aos problemas reprodutivos ocasionados (DUBEY; SCHARES, 2006). O cão doméstico e outros membros da família Canidae são os hospedeiros definitivos do parasita, sendo responsáveis pela eliminação de oocistos no ambiente (MCALLISTER et al., 1998; DUBEY et al., 2011). Exames moleculares, histopatológicos e bioensaio são ferramentas amplamente utilizadas para a identificação dos hospedeiros intermediários, sendo atualmente reconhecido mais de 30 espécies de mamíferos e poucas espécies de aves que atuam como hospedeiro intermediário do *N. caninum* (DUBEY; SCHARES, 2011; DONAHOE et al., 2015).

A infecção nos animais pode ocorrer via vertical, forma mais comum nos bovinos, ou por via horizontal por meio da ingestão de alimentos, água contaminados com oocistos ou ingestão de tecidos de hospedeiros contendo cistos (DUBEY; SCHARES; ORTEGA-MORA, 2007). As aves se infectam pela ingestão de oocistos esporulados presente no ambiente, entretanto, o real papel das aves no ciclo biológico do *N. caninum* ainda é incerto (DONAHOE et al., 2015).

Pombos da espécie *Zenaida auriculata* (Des Murus, 1847) são aves pertencentes à ordem dos Columbiformes, nativas da América Central e do Sul, sendo observadas das Antilhas até a Terra do Fogo (SOUZA et al., 2007). Popularmente conhecidas por nomes como amargosinha, pomba-de-bando e avoante, sua identificação é feita por meio da observação de características como coloração acinzentada, tamanho intermediário de aproximadamente 21 cm, duas faixas negras horizontais na lateral da cabeça e manchas também negras nas asas (SICK, 2001).

Nos últimos anos, a população dessas aves tem aumentado consideravelmente nos estados do Sul do país, principalmente devido ao hábito alimentar dessas aves, que é composta por grãos como trigo, milho, soja, além de semente de plantas e resíduos alimentares produzidos pelo homem (RANVAUD et al., 2001). Em regiões com alta produção desses grãos, essas aves são consideradas como pragas agrícolas, causando elevados prejuízos econômicos

1 (BUCHER; RANVAUD, 2006). Essas aves são consideradas animais sinantrópicos  
2 devido à relação de proximidade com o homem, o que torna preocupante a  
3 transmissão de patógenos, visto que os pombos podem carrear estágios infectantes  
4 de várias doenças, além dos danos ao ambiente público devido ao grande acúmulo  
5 de fezes nos grandes centros urbanos (DIAZ et al., 2008; SHIBATTA et al., 2009;  
6 BARROS et al., 2014).

7 O fato dos pombos serem presas fáceis para os canídeos aliado ao  
8 hábito de alimentação das aves ser realizada principalmente no solo, ficando  
9 expostas à infecção por oocistos presentes no ambiente, torna esses animais como  
10 potenciais hospedeiros intermediário do *N. caninum*, justificando assim estudos que  
11 avaliem a importância epidemiológica dessas aves no ciclo biológico do parasita.

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## 2 REFERENCIAL TEÓRICO

*Neospora caninum* é um parasita intracelular obrigatório pertencente ao filo Apicomplexa. Infecta uma ampla gama de hospedeiros e é causador da neosporose, uma importante doença infecciosa causadora de problemas reprodutivos e síndrome neuromuscular em bovinos e cães, respectivamente, além de já ter sido reportada em outros animais domésticos e silvestres (DUBEY; SCHARES; ORTEGA-MORA, 2007).

O parasita foi descrito pela primeira vez na Noruega por Bjerkås, Mohn e Presthus (1984) que relataram a presença de cistos de protozoários no sistema nervoso central e muscular em cães da raça Boxer, provenientes da mesma ninhada com sinais neurológicos e paresia. Embora a doença descrita fosse semelhante à toxoplasmose, os autores não detectaram anticorpos contra esse parasita, além de observarem uma incapacidade de infectar camundongos. Anos mais tarde, Bjerkås e Presthus (1988) através de estudos imunohistoquímicos e ultraestruturais, concluíram que os cistos observados nesses cães com sinais neurológicos eram diferentes do *T. gondii* e se tratava de uma nova espécie de parasita. No mesmo ano, Dubey et al. (1988) avaliando cortes histológicos de cães que morreram com doença neurológica semelhante à toxoplasmose concluíram a existência de um novo parasita e o denominaram de *N. caninum*.

Após a descrição do parasita até a identificação do seu hospedeiro definitivo, houve um grande intervalo de tempo. Baker et al. (1995) realizaram um estudo experimental com aves carnívoras para avaliar o potencial dessas aves como hospedeiros definitivos do parasita, entretanto os resultados não foram satisfatórios. Somente em 1998, McAllister et al. (1998) descreveram que os cães domésticos (*Canis familiaris*) são hospedeiros definitivos, visto que cães infectados com cistos teciduais eliminaram oocistos não esporulados nas fezes após 8 dias de infecção. Novos estudos foram realizados e mostraram que outros membros da família Canidae, tal qual o coiote (*Canis latrans*) (GONDIM et al., 2004b), o dingo australiano (*Canis lupus dingo*) (KING et al., 2010) e o lobo cinza (*Canis lupus*) (DUBEY et al., 2011) também são hospedeiros definitivos do parasita.

A doença tem sido descrita em vários animais domésticos e silvestres, entretanto bovinos apresentam uma maior susceptibilidade à doença,

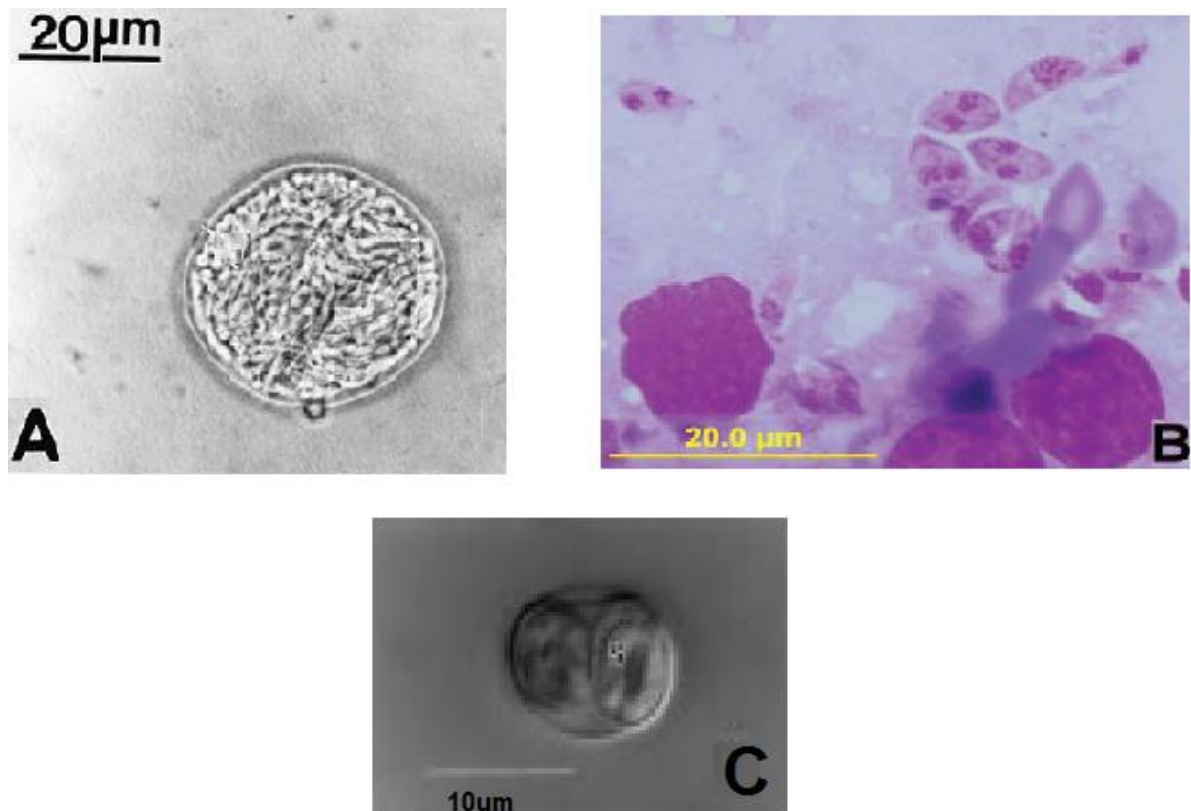
1 devido aos problemas reprodutivos tais como abortamento, natimortalidade e  
2 nascimento de bezerros fracos (DUBEY; SCHARES, 2006, 2011). Estudos  
3 demonstram que a neosporose é reponsável por um prejuízo econômico mundial de  
4 mais de 1,2 bilhões de dólares ao ano, enquanto no Brasil, essas cifras giram em  
5 torno de 153 milhões de dólares, sendo mais de 100 milhões decorrentes da  
6 bovinocultura de corte (REICHEL et al., 2013).

7 Atualmente existem vários testes disponíveis para o diagnóstico  
8 sorológico da neosporose, como o teste de aglutinação (NAT), a imunofluorescência  
9 indireta (RIFI), o ensaio imunoenzimático (ELISA) e o immunoblot (DUBEY;  
10 SCHARES, 2011). Entretanto, a RIFI é considerada como prova padrão para o  
11 sorodiagnóstico devido à baixa reatividade cruzada com outros parasitas coccídios  
12 (BJÖRKMAN; UGGLA, 1999). A presença de anticorpos contra *N. caninum* já foi  
13 observada em felinos (MENESES et al., 2014), camelos (SADREBAZZAZ;  
14 HADDADZADEH; SHAYAN, 2006), galinhas (MARTINS; KWOK; DUBEY, 2011),  
15 equinos (ABREU et al., 2014), lhamas (MORÉ et al., 2008), suínos (FEITOSA et al.,  
16 2014), búfalos (REICHEL et al., 2015), ovinos (ROMANELLI et al., 2007), caprinos  
17 (TOPAZIO et al., 2014) além dos bovinos (SOUSA et al., 2012) e caninos (ROBBE  
18 et al., 2016). Embora a presença de anticorpos anti-*N. caninum* também tenha sido  
19 descrita em seres humanos, o potencial zoonótico ainda é incerto, visto que o  
20 parasita nunca foi identificado ou isolado de tecido humano (TRANAS; HEINZEN;  
21 WEISS, 1999; LOBATO et al., 2006; OSHIRO et al., 2015).

22 Vários animais domésticos já foram descritos como hospedeiros  
23 intermediários do parasita, tais quais os bovinos, ovinos e caprinos (DUBEY et al.,  
24 1996), roedores, animais silvestres como raposa (NASCIMENTO et al., 2015) e  
25 cervídeos (DUBEY; SCHARES, 2011) além de algumas aves, como galinhas  
26 (COSTA et al., 2008), pardal (GONDIM et al., 2010) e corvo (SALANT et al., 2015),  
27 entretanto, o isolamento do parasita somente foi relatado em bovinos (CAMPERO et  
28 al., 2015), búfalos (RODRIGUES et al., 2004), ovinos (KOYAMA et al., 2001), lobo  
29 cinza (DUBEY et al., 2014), veado de cauda branca (VIANNA et al., 2005), bisão  
30 europeu (BIENÍ; MOSKWA; CABAJ, 2010) e cães (DUBEY et al., 1988).

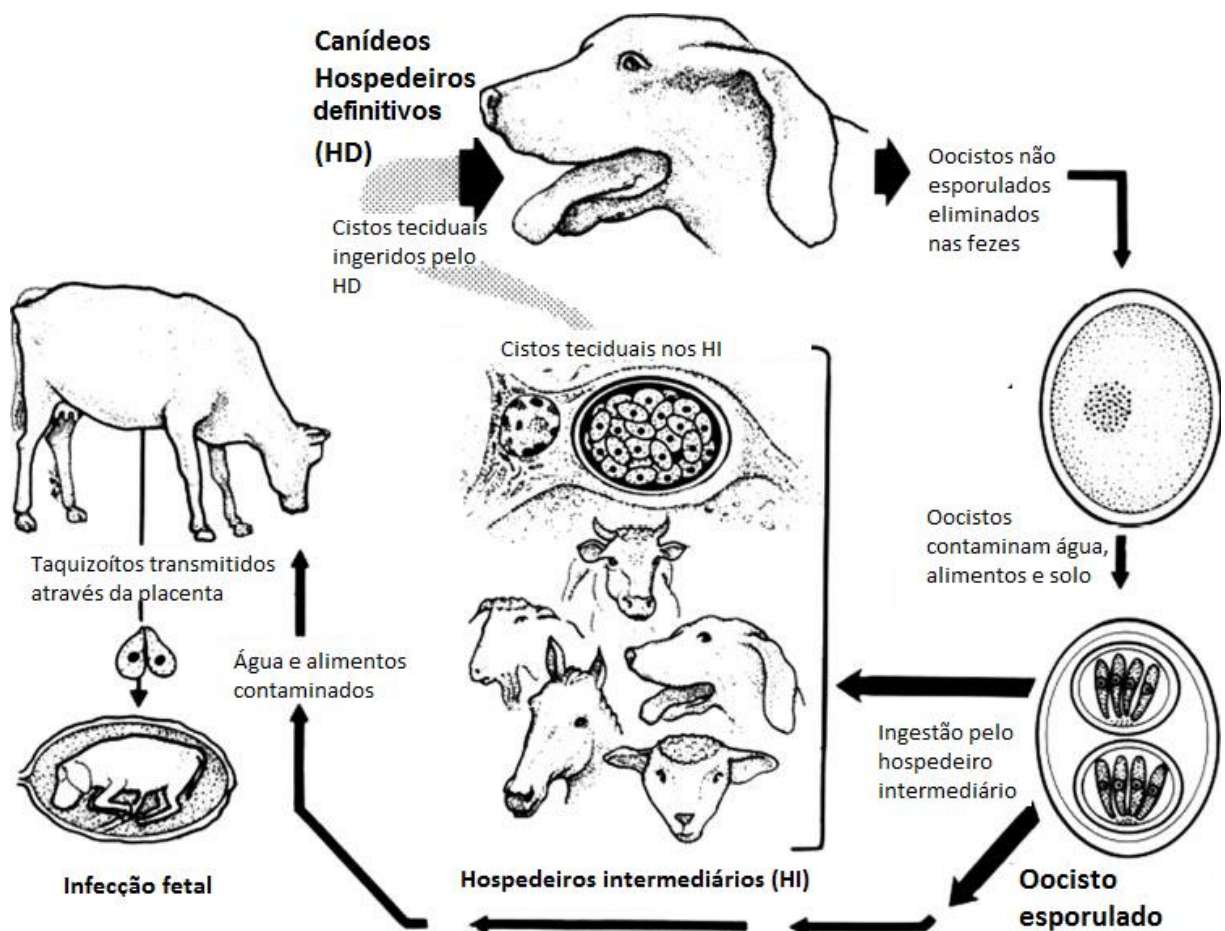
31 O parasita apresenta três formas biológicas distintas: taquizoítos,  
32 bradizoítos e oocistos (Fig.1). Os taquizoítos possuem formato semi-lunar, medem 3-  
33 7µm por 1-5µm e são a forma de multiplicação rápida do parasita, sendo encontrado  
34 no interior de células como hepatócitos, macrófagos alveolares, miócitos, células

1 renais, endoteliais vasculares e nervosas (BARR et al., 1991; DUBEY et al., 2002).  
2 Os bradizoítos são alongados e delgados, medindo aproximadamente 6,5-8 $\mu$ m por  
3 1,5-2 $\mu$ m e envoltos por um cisto. São a forma lenta de multiplicação do parasita,  
4 sendo encontrados principalmente no sistema nervoso central dos hospedeiros  
5 intermediários e definitivos. Possuem formato arredondado ou ovalado e variam no  
6 tamanho, podendo chegar até 107 $\mu$ m de diâmetro e com parede de até 4 $\mu$ m  
7 (DUBEY et al., 2002, 2004). Os oocistos são a forma de resistência do parasita  
8 encontrado no ambiente, medem 10,6 $\mu$ m-12,4 $\mu$ m por 10,6 $\mu$ m-12 $\mu$ m e são  
9 eliminados nas fezes dos canídeos na forma não esporulada. Após esporulação no  
10 meio ambiente, os oocistos são compostos por dois esporocistos, cada qual com  
11 quatro esporozoítos, que são alongados e medem 5,8-7,0 $\mu$ m por 1,8-2,2 $\mu$ m (DUBEY  
12 et al., 2002).



30 **Fig. 1.** Formas parasitárias do *Neospora caninum*. A) Cisto tecidual  
31 com bradizoítos no interior. B) Taquizoítos corados com Giemsa. C) Oocisto  
32 esporulado. Fonte: Adaptado de Lindsay; Upton; Dubey, 1999; Dubey et al., 2004;  
33 Dubey; Schares; Ortega-Mora, 2007

O ciclo biológico do parasita (Fig. 2) é considerado como heteroxeno facultativo, caracterizado pela presença de um canídeo como hospedeiro definitivo, onde irá ocorrer uma ciclo sexual enteroepitelial, ainda não completamente elucidado, culminando com a eliminação de oocistos nas fezes, e de um hospedeiro intermediário, onde ocorrerá uma fase de multiplicação assexuada, culminando com a formação de cistos teciduais (DUBEY; SCHARES, 2006; DUBEY; SCHARES; ORTEGA-MORA, 2007).



**Fig 2.** Ciclo Biológico do *Neospora caninum*. Fonte: Adaptado de Dubey; Schares; Ortega-Mora, 2007.

Os hospedeiros definitivos e intermediários podem se infectar por duas principais vias, horizontal e vertical. A primeira ocorre quando oocistos esporulados são ingeridos com água e alimentos contaminados ou por ingestão de tecidos de hospedeiros intermediários contendo cistos teciduais. A segunda, também

1 chamada de transmissão transplacentária, é bastante comum nos bovinos e ocorre  
2 quando taquizoítos atravessam a barreira placentária e infectam o feto durante a  
3 gestação (DUBEY; SCHARES, 2006; DUBEY; SCHARES; ORTEGA-MORA, 2007).  
4 Vacas prenhes cronicamente infectadas podem ter uma re-agudização da doença,  
5 levando à uma disseminação de taquizoítas e consequente infecção fetal  
6 (WILLIAMS et al., 2009). Transmissão via lactogênea tem sido relatada  
7 experimentalmente bem como DNA do parasita tem sido descrito no sêmen de  
8 touros, entretanto essas duas vias são improváveis de acontecer em condições  
9 naturais (DAVISON et al., 2001; FERRE et al., 2005; DUBEY; SCHARES; ORTEGA-  
10 MORA, 2007).

11 O diagnóstico da neosporose é realizado por meio de exames  
12 sorológicos, já discutidos previamente, ou por meio de técnicas diretas que permitam  
13 a detecção do parasita ou DNA do mesmo, tal como o isolamento, métodos  
14 moleculares e imunohistoquímicos (DUBEY; SCHARES, 2006). O isolamento do  
15 parasita pode ser realizado em animais ou cultivo celular e é de difícil realização  
16 devido à baixa concentração de parasitas em animais naturalmente infectados, além  
17 dos custos elevados e questões bioéticas envolvidas (DUBEY; SCHARES;  
18 ORTEGA-MORA, 2007). O exame histopatológico deve ser utilizado como método  
19 auxiliar no diagnóstico pois apresenta baixa sensibilidade de detecção do parasita  
20 em cortes de tecidos, devendo ser sempre aliado com a imunohistoquímica, devido à  
21 utilização de anticorpos poli ou monoclonais para a identificação específica de  
22 antígenos parasitário (DUBEY; SCHARES, 2006).

23 Métodos moleculares como a PCR convencional e nested-PCR são  
24 técnicas de elevada sensibilidade e especificidade, sendo aplicáveis à uma ampla  
25 variedade de amostras biológicas. O gene Nc5 e a região do espaçador transcrito  
26 interno 1 (ITS-1) do RNA são os marcadores mais comumente utilizados para  
27 detecção do parasita, sendo o ITS-1 também utilizado para análises filogenéticas,  
28 pois apresenta regiões conservadas intraespécie e variáveis interespecies,  
29 permitindo assim a diferenciação entre outros coccídios (GONDIM et al., 2004a;  
30 DUBEY; SCHARES, 2006; DONAHOE et al., 2015). Outras técnicas moleculares  
31 como o mini e microsátélites estão disponíveis para identificação da variabilidade  
32 genética, o que pode determinar em uma variação na resposta imunológica ao  
33 parasita com consequente diferenças na patogenicidade da infecção (REGIDOR-  
34 CERRILLO et al., 2013).

1                   Medidas de controle devem ser tomadas para evitar os prejuízos  
2 econômicos causados pela neosporose, principalmente em bovinos. Dentre essas  
3 medidas, são descritas a realização de exames sorológicos antes da introdução de  
4 animal no rebanho, evitar o acesso de cães à áreas de armazenamento de comida  
5 dos animais, descarte adequado de fetos abortados e restos placentários, prevenção  
6 da contaminação da água com fezes de canídeos, controle de roedores,  
7 transferência de embrião e abate de animais infectados (DUBEY; SCHARES;  
8 ORTEGA-MORA, 2007).

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**3 ARTIGO DE REVISÃO PARA PUBLICAÇÃO\*\***

**NEOSPORA CANINUM IN BIRDS: A REVIEW**

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## 1 **Abstract**

2 *Neospora caninum* is an obligate intracellular protozoan parasite that infects  
3 domestic and wild animals. Canids are considered definitive hosts since they may  
4 shed oocysts in the environment by feces. The disease is recognized as one of the  
5 major causes of bovine abortion worldwide, leading to important economic losses in  
6 dairy and beef cattle. Previously studies have been reported *N. caninum* infection in  
7 different species of birds, what have been associated with increase in seroprevalence  
8 and reproductive problems in dairy cattle. Although the role of birds in the  
9 epidemiological cycle of neosporosis is unknown, these animals are exposed to  
10 infection, since they feed on the ground, and could contribute to parasite  
11 dissemination. This review is focused on current status of neosporosis in birds.

12

13 **Keywords:** *Neospora caninum*; *neosporosis*; *birds*

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## 15 **1. Introduction**

## 16 **2. Serological studies**

## 17 **3. Experimental infections**

## 18 **4. Natural infections**

## 19 **5. Conclusions**

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## 21 **1. Introduction**

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23 *Neospora caninum* is a protozoan parasite that was first described by Bjerkås  
24 et al. (1984), who had reported dogs with cysts-forming sporozoon associated with  
25 lesions in the central nervous system and skeletal muscles. Since it shares

1 morphological similarity with *T. gondii*, *N. caninum* has been misdiagnosed until 1988  
2 however Bjerkås and Presthus (1988) showed by immuno-histochemical and  
3 ultrastructural characteristics of cysts, they were different. Due to these discoveries,  
4 *N. caninum* was formally recognized as a new genus and species (Dubey et al.,  
5 1988a).

6 The life cycle of *N. caninum* was unclear at the beginning and an  
7 experimental study with carnivorous birds was performed to determine if birds were  
8 definitive host of *N. caninum* (Baker et al., 1995). This study used the red-tailed  
9 hawks (*Buteo jamaicensis*), turkey vultures (*Cathartes aura*), barn owls (*Tyto alba*)  
10 and the american crows (*Corvus brachyrhynchus*), that were fed with mice that had  
11 been inoculated subcutaneously with  $10^5$  culture-derived tachyzoites of *N. caninum*.  
12 Although occasional unsporulated oocysts were observed in fecal samples, none of  
13 these were similar to *N. caninum* and mice inoculated with these oocysts did not  
14 show either antibodies against *N. caninum* or tissue cysts in the brain.

15 A few years later, McAllister et al. (1998) showed that dogs (*Canis familiaris*)  
16 are definitive hosts of the parasite, since a sexual phase occurs in the intestine of  
17 these animals shedding oocysts in their feces. After the description of dogs as  
18 definitive hosts of the *N. caninum* life cycle, other studies showed that canids,  
19 including the Australian dingo (*Canis lupus dingo*) (King et al., 2010), the coyote  
20 (*Canis latrans*) (Gondim et al., 2004) and the gray wolf (*Canis lupus*) (Dubey et al.,  
21 2011) are definitive hosts and can shed oocysts in their feces. Afterwards, Dubey et  
22 al. (2002) based mainly on morphologic aspects, made a redescription of the parasite  
23 and a differentiation of related parasites.

24 In the past years, many studies using molecular techniques have been used to  
25 show that small mammals and birds are intermediate hosts of *N. caninum*

1 (Čobádiová et al., 2013; Costa et al., 2008; Du et al., 2015; Ferroglio et al., 2007;  
2 Fuehrer et al., 2010; Gondim et al., 2010; Hughes et al., 2008; Truppel et al., 2010).  
3 However, viable parasite has been isolated only in dog (Dubey et al., 1998), gray wolf  
4 (Dubey et al., 2014), cattle (Campero et al., 2015), sheep (Pena et al., 2007), water  
5 buffalo (Rodrigues et al., 2004), bison (Bień et al., 2010) and white-tailed deer  
6 (Vianna et al., 2005). Antibodies against this parasite has been described in many  
7 hosts, even in human, however the zoonotic potential is uncertain since the parasite  
8 has not been detected in human tissues (Tranas et al., 1999)

9         Although the role of birds in the life cycle of *N. caninum* is not completely  
10 understood, previously studies have shown that the presence of birds on dairy farms  
11 could increase the seroprevalence and reproductive problems associated to *N.*  
12 *caninum*, suggesting that these animals may be an important intermediate host,  
13 contributing to transmission of the parasite to definitive hosts (Bartels et al., 1999;  
14 Otranto et al., 2003). Thus, the present manuscript provides an overview on what is  
15 known about *N. caninum* in birds.

16

## 17 **2. Serological studies**

18

19         There are few serological studies to detect anti-*N. caninum* antibodies in birds.  
20 The first one was performed by Costa et al. (2008), which using indirect fluorescent  
21 antibody tests (IFAT) observed a seroprevalence of 12.5 % (50/400) in chickens from  
22 Bahia state, Brazil. However, when compared the origin of these animals, the  
23 prevalence was higher in outdoors chickens (23.5%, 47/200) than indoor (1.5%,  
24 3/200), showing that chickens raised outside are more exposed to *N. caninum* and  
25 may contribute to the parasite transmission. Gonçalves et al. (2012) also working

1 with chickens from Brazil have observed a prevalence of 17% (17/100) using IFAT  
2 and a cut off of 50, moreover, an significant agreement between *T. gondii* and *N.*  
3 *caninum* serology was observed, indicating that birds can may have contact with  
4 both coccidian parasites.

5         Afterwards, a serological study with wild and captive birds was performed  
6 (Mineo et al., 2011). The authors used 294 birds, belonging to nine different orders,  
7 but no serological positivity was observed in all tested samples. Although they did not  
8 observed antibodies against *N. caninum* in sera from these animals, Apicomplexa-  
9 like cysts immunostained with *N. caninum* antisera were found in cloacae and  
10 cervical musculature from two Psittaciformes.

11         In the same year, Martins et al. (2011) performed a serological study of *N.*  
12 *caninum* in free-range chickens (*Gallus domesticus*) from the Americas. Birds from  
13 previous studies with *T. gondii* from different countries of North, Central and South  
14 America were used. According with IFAT and a cut off of 25, antibodies against the  
15 parasite was found in 39.5% (524/1324) of birds, showing that free-range chickens  
16 have been exposed to *N. caninum*.

17         Molina-López et al. (2012) observed a high prevalence of antibodies against  
18 *N. caninum* in common raven (*Corvus corax*) from Northeast of Spain. Using IFAT and  
19 a cut off of 50, 35.8% (24/67) of animals were positive, with titers ranging from 50  
20 until 100. A recent study with free ranging crows was done in Israel (Salant et al.,  
21 2015). A total of 183 birds belonging to the species *Corvus cornix*, *Corvus monedula*  
22 and *Corvus splendens* were used to evaluate the exposure of these animals to *N.*  
23 *caninum* by means of modified agglutination test (MAT) and IFAT. Thirty out 183  
24 animals (16.4%) were positive in both tests, showing that crows (*C. cornix* and *C.*

1 *monedula*) are exposed to *N. caninum* and could be an important role in the  
2 epidemiology of the parasite.

3       Recently, Camillo et al. (2015) performed a study with backyard chickens,  
4 since they may be good bioindicator of the environmental contamination with oocysts.  
5 One hundred and thirty-seven animals from 23 different farms from Rio Grande do  
6 Sul state, Brazil were tested using IFAT and cut off of 50. Antibodies anti-*N. caninum*  
7 were detected in 36.5% (50/137), indicating a widespread exposure of backyard  
8 chickens to the parasite in that region.

9       A summary of seroprevalence studies of *N. caninum* in birds is shown in Table  
10 1.

11

### 12 **3. Experimental infections**

13

14       The first experimental study with *N. caninum* and birds was performed by  
15 Baker et al. (1995) when definitive host of the parasite was unknown. After the  
16 discovery that canids are definitive hosts, some experimental studies were done to  
17 evaluate the potential of birds as intermediate hosts in the life cycle. In the past, only  
18 cattle were known as intermediate host of *N. caninum*, but since serological studies  
19 have provided evidence of infection in wild canids, McGuirre et al. (1999) proposed  
20 that *N. caninum* would have an intermediate host that is common pray of those wild  
21 canids. Thus, the authors designed an experimental study with inoculation of *N.*  
22 *caninum* tachyzoites in pigeons (*Columba livia*) and zebra finches (*Poephila guttata*).  
23 Three animals from each species were inoculated intraperitoneally with different  
24 doses ( $10^6$ ,  $10^5$  and  $10^4$ ) of two different strains (NC-2 and NC-Liverpool) of the  
25 parasite. By means of IFAT, tissue culture and PCR, all pigeons were positive and

1 none of zebra finches were. Although only one pigeon was positive in the histological  
2 examination, typical lesions of neosporosis were observed. These results have  
3 suggested that Columbiform birds could be a good candidate for intermediate host  
4 and could play an important role in the biological cycle.

5 Furuta et al. (2007) realized an experimental infection in chickens and  
6 embryonated eggs with *N. caninum*. Laying hens, 7-day-old chicks and embryonated  
7 eggs were inoculated with different parasite concentration. None of chicks showed  
8 clinical signs of infection but they had seroconversion, with titer of 400 on IFAT 15  
9 days post-infection (d.p.i.). Immunohistochemistry analysis also showed the parasite  
10 in different organs, showing disseminated acute infection. Laying hens showed a  
11 similar pattern to chickens in serological response, with positive result on IFAT 15  
12 d.p.i. and non-detectable antibodies at 60 d.p.i. None evidence of vertical  
13 transmission was observed, but embryonated eggs infected by through allantoic  
14 cavity had susceptibility to infection with high mortality rate. Dogs that had received  
15 inoculated eggs produced serological response and shed oocysts by feces, showing  
16 that chickens could be an intermediate host of *N. caninum*.

17 In the same line of research, an experimental study with broiler chicken  
18 embryonated eggs was conducted (Mansourian et al., 2009). Embryonated eggs  
19 were inoculated with different tachyzoites concentrations (from 10 to 10<sup>6</sup>) via  
20 chorioallantoic. None hatching was observed in the groups with the highest parasite  
21 concentration, while in the groups inoculated with low parasite concentration,  
22 hatching occurred, but one chicken showed neurological signs such as hind limb  
23 paralysis, pedaling movements, lack of coordination, while other three birds had  
24 arthritis in the feet joints.

1           The potential of quails (*Coturnix coturnix japonica*) as an intermediate host of  
2 *N. caninum* was also evaluated (de Oliveira et al., 2013). Animals were divided in  
3 three groups and received different concentrations of parasite ( $3.5 \times 10^6$ ,  $5 \times 10^6$  and  
4 placebo group) subcutaneously. None of the quails showed clinical signs of infection  
5 or died during the experiment. By means of IFAT and a cut off of 10, seroconversion  
6 occurred on 7 d.p.i., with peak on 14 d.p.i. Mononuclear inflammatory infiltrate,  
7 immunoreactivity by IHC and molecular detection was observed mainly during the  
8 first two weeks of infection. None oocyst was found in feces from dogs that were fed  
9 with inoculated quails, no molecular detection of dog stool samples and no dog  
10 seroconversion was observed, demonstrating that quails are not a good intermediate  
11 host for *N. caninum*.

12           Munhoz et al. (2014) analyzed the susceptibility of *Gallus gallus domesticus* to  
13 *N. caninum* tachyzoites infection. For this, two experiments were done. In the first  
14 one, poultries were divided in groups that received feed with and without coccidiostat,  
15 followed by subcutaneous inoculation of  $3 \times 10^6$  tachyzoites. No evidence of infection  
16 in adults and chicks were observed by IHC and histopathology. It also was not  
17 observed DNA amplification in infected bird tissues and no dogs that were fed by  
18 tissues from birds shed oocyst by feces. In another experiment, embryonated chicken  
19 eggs were challenged with  $1 \times 10^2$  tachyzoites. Chicks born from these eggs were  
20 euthanized and again, no lesion compatible with neosporosis, no DNA amplification  
21 and no oocysts shed by dogs were observed, indicating that the parasite could have  
22 been eliminated by the host.

23

#### 24 **4. Natural infections**

25

1           Although antibodies against *N. caninum* have been described in many  
2 animals, such as canids, felids, cervids, ruminants, birds and marine mammals, just  
3 few of these are considered as intermediate host. The first study that showed that  
4 birds are intermediate host for *N. caninum* was done by Costa et al. (2008). Brain  
5 tissue from 10 positive chickens by serological test was used to DNA extraction  
6 followed by PCR. Cloning of PCR fragments and sequencing were also performed.  
7 Six animals (60%) from 10 seropositive were positive by PCR for *N. caninum* and  
8 negative for *T. gondii*, showing for the first time a natural *N. caninum* infection in  
9 birds. In another study with natural infection of chickens from Brazil, DNA from *N.*  
10 *caninum* was detected in 6% of animals, using brain and heart tissue (Gonçalves et  
11 al., 2012). By means of nested-PCR based on ITS-1 region followed by sequencing,  
12 the sequences showed 99-100% of identity with previously data. Furthermore, all  
13 animals positive by PCR were negative at serological tests, showing that birds can be  
14 infected but seronegative, probably due to decrease of specific circulating antibodies  
15 in infected birds as previously demonstrated by Mineo et al. (2009).

16           Afterwards, Gondim et al. (2010) showed *N. caninum* infection in sparrows  
17 (*Passer domesticus*) from Brazil. Heart and brain of 40 animals from Pernambuco  
18 and Bahia states were used for DNA extraction followed by a nested PCR using  
19 primers to conserved region of internal transcribe spacer (ITS1) between *Hammondia*  
20 sp., *N. caninum* and *T. gondii*. Sequentially, sequencing was performed to identify the  
21 parasite. In the nested-PCR, 25% (10/40) of samples were positive for  
22 Toxoplasmatinae, however, 30% of those positive samples were classified as *N.*  
23 *caninum* by DNA sequencing. It was the first study that showed that wild birds may  
24 act as intermediate host of *N. caninum*, indicating an importance of these birds in  
25 epidemiological cycle, since sparrows, may be prey by canids, contributing to

1 transmission of the parasite. Abdoli et al. (2015) have also detected *N. caninum* in  
2 sparrows from Iran by molecular tests. Brain tissue from 217 birds were used for  
3 DNA extraction and nested-PCR using primers Np21 plus and Np6plus in the first  
4 reaction and Np7 and Np6 in the second. DNA from *N. caninum* was observed in  
5 eight birds (3.68%) and sequence analyses showed 97-99% of similarity with other  
6 isolates deposited in GenBank.

7         The presence of tissue cysts in wild birds was previously described (Mineo et  
8 al., 2011). Analysis of two birds, one red-and-green macaw (*Ara chloropterus*) and  
9 other blue-fronted Amazon parrots (*Amazona aestiva*), that had died during  
10 internment in a veterinary hospital, showed Apicomplexan-like tissue cysts in HE  
11 stained from the cloacae of the first bird and cervical of the second.  
12 Immunohistochemistry analysis of these samples revealed positive immunostaining  
13 with anti-*N. caninum* polyclonal antibodies. This first report of latent parasitic forms in  
14 the musculature of wild birds brought some important points to discussion of  
15 *Neospora* life cycle in wildlife animals.

16         *Neospora caninum* DNA has also been detected in brain tissue from wild birds  
17 of Spain (Darwich et al., 2012). Brain tissue from 201 animals belonging to 14  
18 species from three different regions of Spain was used for DNA extraction and PCR  
19 test using the primers Np21-plus and Np6-plus, targeting genomic Nc5 region. Only  
20 three (1.5%) samples were positive for *N. caninum* DNA, two from magpies (*Pica*  
21 *pica*) and one from common buzzard (*Buteo buteo*), showing 98% of identify with  
22 previously *N. caninum* sequences described and deposited in GenBank. The  
23 detection of *N. caninum* in these birds should be considered important for the  
24 epidemiologic cycle, since common buzzard is a common predator and usually ingest  
25 carrion, which could contribute to infection of these animals.

1           A study with crows of three different species from different areas of Israel was  
2 conducted to confirm if these animals were an intermediate host of *N. caninum*  
3 (Salant et al., 2015). Nested-PCR targeting Nc5 gene was carried out to detect the  
4 DNA of the parasite in the brain tissue from crows. Only two animals (1.09%)  
5 belonging to the species *Corvus cornix* and *Corvus monedula*, were found to be  
6 positive by nPCR, showing 99% of identify with DNA sequences previously  
7 described. Interesting, both positive samples were from animals seronegative by  
8 means of MAT and IFAT assay, which point for the importance of the molecular test  
9 since some birds may be present an undetectable level of antibodies after infection  
10 with *N. caninum*.

11           Recently, DNA of the parasite was described in pigeons from China (Du et al.,  
12 2015) Brain tissue of 210 pigeons from pigeons farms were collected for DNA  
13 extraction. DNA detection of parasite was performed by nested-PCR using primers  
14 for specific Nc5 gene while ITS-1 region were used to sequencing analysis. PCR  
15 positive result was observed in 63 (30%) and ITS-1 analysis showed the presence of  
16 clusters with 42 samples assigned to the NC-PR genotype, 10 to NC-1 genotype and  
17 5, 3 and 3 to new genotypes identified. This was the first report of *N. caninum* DNA in  
18 tissues from pigeons, which expands the list of the intermediate host of *N. caninum*  
19 and point for the importance of these animals in the transmission of the parasite.

20           A summary of all birds considered intermediate hosts of *N. caninum* is shown  
21 in Table 2 while other species in Table 3.

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## 5. Conclusions

Nowadays, molecular studies have shown that *N. caninum* has a large range of intermediate host, which contributes to elucidate the epidemiology of the disease. Birds in special may have contribute to dissemination of the parasite, mainly because these animals usually feed on the ground, becoming expose to many pathogens, and are preyed upon by canids, contributing to the life cycle of the parasite.

Further studies with different species of birds are needed to elucidate the real importance of birds in the epidemiology of neosporosis, aiming control the disease and decrease the economic impact.

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23

**Table 1.** Serological prevalence of *Neospora caninum* in birds from different countries.

Animal specie	Country	Seroprevalence rate (%)	Test	Cut off	Reference
Chicken ( <i>Gallus domesticus</i> )	Brazil	12.5	IFAT	1:50	Costa et al., 2008
Chicken ( <i>Gallus domesticus</i> )	Mexico	18.5	IFAT	1:25	Martins et al., 2011
	USA	7.2			
	Costa Rica	39.5			
	Grenada	71.5			
	Guatemala	44.0			
	Nicaragua	83.6			
	Argentina	58.1			
	Brazil	34.3			
	Chile	62.3			
	Colombia	11.2			
	Guyana	38.7			
	Peru	18.0			
	Venezuela	21.7			
Chicken ( <i>Gallus domesticus</i> )	Brazil	17	IFAT	1:50	Gonçalves et al., 2012
Chicken ( <i>Gallus domesticus</i> )	Brazil	36.5	IFAT	1:50	Camillo et al., 2015
Black vulture ( <i>Coragyps atratus</i> )	Brazil	0	IFAT	1:20	Mineo et al., 2011
Rock dove ( <i>Columba livia</i> )	Brazil	0			
Eared dove ( <i>Zenaida auriculata</i> )	Brazil	0			
Southern caracara ( <i>Caracara plancus</i> )	Brazil	0			
Great-billed seed-finch ( <i>Oryzoborus maximiliani</i> )	Brazil	0			
Atlantic canary ( <i>Serinus canaria</i> )	Brazil	0			
Toco toucan ( <i>Ramphastos toco</i> )	Brazil	0			
Blue fronted Amazon parrot ( <i>Amazona aestiva</i> )	Brazil	0			
Hyacinth macaw ( <i>Anodorhynchus hyacinthinus</i> )	Brazil	0			
Lear's macaw ( <i>Anodorhynchus leari</i> )	Brazil	0			
Blue-and-yellow macaw ( <i>Ara ararauna</i> )	Brazil	0			
Red-and-green macaw ( <i>Ara chloropterus</i> )	Brazil	0			
Budgerigar ( <i>Melopsittacus undulatus</i> )	Brazil	0			

Striped owl ( <i>Asio clamator</i> )	Brazil	0			
Barn owl ( <i>Tyto alba</i> )	Brazil	0			
Greater rhea ( <i>Rhea americana</i> )	Brazil	0			
Common ostrich ( <i>Struthio camelus</i> )	Brazil	0			
Common raven ( <i>Corvus corax</i> )	Spain	35.8	IFAT	1:50	Molina-López et al., 2011
Hooded crow ( <i>Corvus cornix</i> )	Israel	17.9	MAT/IFAT	1:100/1:50	Salant et al., 2015
Western jackdaw ( <i>Corvus monedula</i> )	Israel	20.0			
House crow ( <i>Corvus splendens</i> )	Israel	0			

**Table 2.** Birds as natural intermediate hosts for *Neospora caninum* due to detection of the parasite using different methods from bird tissues.

Animal specie	Location	Method	Tissue	Reference
Blue fronted Amazon parrot ( <i>Amazona aestiva</i> )	Brazil	IHC	Cervical musculature	Mineo et al., 2011
Red-and-green macaw ( <i>Ara chloropterus</i> )	Brazil	IHC	Cloacae musculature	Mineo et al., 2011
Chickens ( <i>Gallus domesticus</i> )	Brazil	PCR (Nc5)	Brain	Costa et al., 2008
Sparrow ( <i>Passer domesticus</i> )	Brazil	PCR (ITS-1)	Brain/Heart	Gondim et al., 2010
Hooded crow ( <i>Corvus cornix</i> )	Israel	PCR (Nc5)	Brain	Salant et al., 2015
Western jackdaw ( <i>Corvus monedula</i> )	Israel	PCR (Nc5)	Brain	Salant et al., 2015
Magpies ( <i>Pica pica</i> )	Spain	PCR (Nc5)	Brain	Darwich et al., 2012
Common buzzard ( <i>Buteo buteo</i> )	Spain	PCR (Nc5)	Brain	Darwich et al., 2012

**Table 3.** Mammals as natural intermediate hosts for *Neospora caninum* due to detection of the parasite using different methods from animal tissues.

<b>Animal species</b>	<b>Location</b>	<b>Method</b>	<b>Tissue</b>	<b>Reference</b>
Cattle ( <i>Bos taurus</i> )	Argentina	PCR (Nc5), mouse bioassay	Brain	Campero et al., 2015.
European bison ( <i>Bison bonasus bonasus</i> )	Poland	Culture isolation	Blood	Bi�n et al., 2010
Field mouse ( <i>Apodemus sylvaticus</i> )	Italy	PCR (Nc5)	Kidney and skeletal muscle	Ferroglio et al., 2007.
House mice ( <i>Mus musculus</i> )	UK	PCR (Nc5)	Brain	Hughes et al., 2006.
Rats ( <i>Rattus norvegicus</i> )	Taiwan	PCR (Nc5)	Brains	Huang et al., 2004.
Sheep ( <i>Ovis aries</i> )	Brazil	Dogs bioassay/PCR (Nc5)	Brain	Pena et al., 2007.
White-tailed deer ( <i>Odocoileus virginianus</i> )	USA	Mouse bioassay	Brain	Vianna et al., 2005.
Water buffaloes ( <i>Bubalus bubalis</i> )	Brazil	Dogs/gerbils bioassay	Brain	Rodrigues et al., 2004.
Dogs ( <i>Canis familiaris</i> )	Brazil	Gerbil bioassay	Brain	Gondim et al., 2001.
Capybaras ( <i>Hydrochaeris hydrochaeris</i> )	Brazil	PCR (Nc5/ITS-1)	Lymph nodes, heart, liver and blood	Truppel et al., 2010.
Goat ( <i>Capra hircus</i> )	Argentina	Mouse/gerbils bioassay	Fetal fluid and brain, placenta	Unzaga et al., 2014.
Hoary fox ( <i>Pseudolopex vetulus</i> )	Brazil	PCR (Nc5)	Brain	Nascimento et al., 2015.
Axis fawn ( <i>Axis axis</i> )	Argentina	PCR (Nc5) and mouse/gerbil bioassay	Brain	Basso et al., 2014.
European brown bear ( <i>Ursus arctos</i> )	Slovakia	PCR (Nc5/ITS-1)	Liver and spleen	Bartley et al., 2013
Mink ( <i>Neovision vision</i> )	UK	PCR (ITS-1)	Brain	Bartley et al., 2013
Fox ( <i>Vulpes vulpes</i> )	UK	PCR (ITS-1)	Brain	Bartley et al., 2013
Polecat ( <i>Mustela putorius</i> )	UK	PCR (ITS-1)	Brain	Bartley et al., 2013
Ferret ( <i>Mustela furo</i> )	UK	PCR (ITS-1)	Brain	Bartley et al., 2013
Badger ( <i>Meles meles</i> )	UK	PCR (ITS-1)	Brain, neck muscle, liver	Bartley et al., 2013

Fallow Deer ( <i>Dama dama</i> )	Switzerland	IHC, HP, PCR (Nc5)	Spinal cord, brain	Soldati et al., 2004.
Blue foxes ( <i>Alopex lagopus</i> )	China	PCR (Nc5), HP, IHC	Brain	Yu et al., 2009.
Red foxes ( <i>Vulpes vulpes</i> )	Czech Republic	PCR (Nc5)	Brain	Hůrková, Modrý, 2009.
Lesser kudu ( <i>Tragelaphus imberbis</i> )	Germany	PCR (Nc5)	Brain, lung, heart, liver and spleen	Peters et al., 2001.
Raccoon ( <i>Procyon lotor</i> )	USA	IHC, PCR (Nc5)	Brain	Lemberger et al., 2005
Black-tailed fawn ( <i>Odocoileus hemionus columbianus</i> )	USA	HP, IHC	Kidney, liver and lung	Woods et al., 1994.
Eld's deer ( <i>Cervus eldi siamensis</i> )	France	IHC	Brain	Dubey et al., 1996.
Alpaca ( <i>Vicugna pacos</i> )	Peru	IHC, PCR (ITS-1)	Brain	Serrano-Martínez et al., 2004.
Llamas ( <i>Llama glama</i> )	Peru	IHC, PCR (ITS-1)	Brain	Serrano-Martínez et al., 2004.
White rhinoceros ( <i>Cerathotherium simum</i> )	South Africa	IHC	Heart	Williams et al., 2002.
Vole ( <i>Microtus arvalis</i> )	Austria	PCR (Nc5)	Brain	Fuehrer et al., 2010
Water vole ( <i>Arvicola terrestris</i> )	Austria	PCR (Nc5)	Brain	Fuehrer et al., 2010
Wild rabbits ( <i>Oryctolagus cuniculus</i> )	UK	PCR (Nc5)	Brain, heart and tongue	Hughes et al., 2008
Rock squirrel ( <i>Spermophilus variegatus</i> )	Mexico	HP, IHC, PCR (ITS-1)	Liver, heart, spinal cord and brain	Medina-Esparza et al., 2013.
European Pine Marten ( <i>Martes martes</i> )	Netherlands	HP, IHC	Hert, liver and lung	Van der Hage et al., 2002
Parma wallaby ( <i>Macropus parma</i> )	Austria	HP, IHC, PCR (ITS-1)	Heart	Cronstedt-Fell et al., 2012

## 4 OBJETIVOS

### 4.1 OBJETIVO GERAL

- Avaliar a importância dos pombos (*Zenaida auriculata*) na epidemiologia do *Neospora caninum*.

### 4.1 OBJETIVOS ESPECÍFICOS

- Padronizar uma técnica de ELISA para a pesquisa de anticorpos anti-*N. caninum* em pombos *Z. auriculata*.
- Verificar a ocorrência de *N. caninum* em pombos (*Z. auriculata*) naturalmente infectados capturados no município de Londrina, Paraná.
- Avaliar por meio da infecção experimental o potencial dos pombos (*Z. auriculata*) de atuarem como hospedeiro intermediário do *N. caninum*.

**5 ARTIGO PARA PUBLICAÇÃO I\*\***

**SURVEY OF *NEOSPORA CANINUM* IN EARED DOVES (*ZENAIDA AURICULATA*) IN SOUTHERN BRAZIL**

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## 1 **Abstract**

2 Neosporosis is an infectious disease caused by *Neospora caninum*, a protozoan  
3 parasite that has worldwide distribution and is responsible for enormous economic  
4 losses in cattle. Birds are considered a good bioindicator of environmental  
5 contamination, since they feed on the ground, being exposed to *N. caninum* oocysts.  
6 The aim of this study was to determine the occurrence of antibodies against *N.*  
7 *caninum* and to verify the presence parasite DNA in tissue from free-ranging eared  
8 doves (*Zenaida auriculata*) from Southern Brazil. For this purpose, serum and brain  
9 samples were collected from 249 doves for ELISA and PCR analysis respectively.  
10 The prevalence of *N. caninum* antibodies in doves was 31.72% (79/249) and  
11 detection of DNA was not observed in none of birds. This is the first report of  
12 antibodies against *N. caninum* in doves *Z. auriculata*, what show us that these birds  
13 had previously contact with the parasite but since no DNA was detected, more  
14 studies should be performed to elucidate the real importance of doves in the  
15 epidemiologic cycle of the *N. caninum*.

16

17 **Keywords:** *Neospora caninum*; *Zenaida auriculata*; serology; ELISA; PCR

18

## 19 **1. Introduction**

20

21 *Neospora caninum* is a coccidian parasite of animals that mainly affect cattle,  
22 being considered as one of the major responsible for reproductive problems by  
23 infectious agents in cattle worldwide (Reichel et al., 2013). Canids, including dogs  
24 (*Canis familiaris*), Australian dingoes (*Canis lupus dingo*), coyotes (*Canis latrans*)  
25 and gray wolf (*Canis lupus*) are definitive hosts for this parasite, however it can infect

1 different species of mammals and birds (Darwich et al., 2012; Dubey et al., 2011;  
2 Gondim et al., 2004; King et al., 2010; McAllister et al., 1998). An association  
3 between the presence of birds on dairy farms and reproductive problems associated  
4 with *N. caninum* has been reported, suggesting that these animals could play an  
5 important role in the epidemiology of this disease (Bartels et al., 1999; Ould-  
6 Amrouche et al., 1999).

7 Birds of *Zenaida auriculata* species (Des Murus, 1847), commonly known as  
8 eared doves, belong to Columbiformes order and occur from the Antilles to Tierra del  
9 Fuego. These animals are considered sinantropic, since they are not domesticated  
10 animals and live close to humans, mainly due to the high food supply, favorable  
11 environmental conditions and absence of predator, leading to population growth out  
12 of control, generating consequently agricultural losses, aesthetic damage to buildings  
13 and problems on public health. Moreover, these birds are an important source of food  
14 for humans in some regions of Brazil (Souza et al., 2007) and may contribute to  
15 transmission of zoonotic diseases (Cano-Terriza et al., 2015; Cong et al., 2012; Li et  
16 al., 2015)

17 The aim of the present work was to study *N. caninum* infection in free-ranging  
18 eared doves from Southern Brazil.

19

## 20 **2. Material and methods**

21

### 22 2.1 Animals and study locations

23 Two hundred and forty-nine eared doves (*Z. auriculata*), including 114 males  
24 and 135 females were trap captured between January 2010 and December 2011 in  
25 three different region of Londrina (23°08'47" to 23°55'46"S/50°52'23" to 51°19'11"W)

1 Paraná state, Southern Brazil. Those animals were obtained from soybean seed  
2 plant (n= 140), campus of the State University of Londrina (n= 80) and a dairy cattle  
3 farm (n= 29). All experimental procedures were approved by Animal Ethics  
4 Committee of State University of Londrina (CEEA no. 70/08) and the Brazilian  
5 Institute of the Environment (IBAMA-SISBIO no. 16.428-1)

6

## 7 2.2 Sample collection

8 All doves were euthanized following the guidelines established by Nacional  
9 Council for Animal Experimental Control (CONCEA/Brazil). Blood collection was  
10 performed by cardiac puncture and the resultant serum samples were stored at -  
11 20°C until further analysis. Brain tissue from doves were collected aseptically and  
12 submitted to molecular detection of the parasite.

13

## 14 2.3 Serology

15 In order to detect antibodies IgG against *N. caninum*, an enzyme-linked  
16 immunosorbent assay test (ELISA) was developed based on previously studies (Paré  
17 et al., 1995). For the preparation of antigen, tachyzoites of NC-1 strain were growth in  
18 Vero cell supplemented with RPMI medium, bovine fetal serum and penicillin (10,000  
19 U) and streptomycin (10mg/ml). Subsequently, the cells were scraped and passed  
20 through 27G needles. The resultant solution was centrifuged 10,000 g for 10 min and  
21 the pellet was mixed with glass beads through vortex three times of 10 minutes each.  
22 The solution was centrifuged again following the same conditions described before  
23 and the pellet was discarded. Antigen solution was submitted to protein concentration  
24 measure by using a commercial kit (Pierce™ BCA, Thermo Fisher, USA). The  
25 antigen was stored at -20° until performance of serological tests.

1           Previously tests were conducted to determine the optimal antigen  
2 concentration and dilutions of serum and conjugate. Flat-bottom 96 well polystyrene  
3 microtitration plates (Nunc-Immuno Plate, MaxiSorp, Denmark) were coated with  
4 0.1ml of crude antigen (2.5µg/ml) diluted in 0.1M carbonate buffer (pH 9.6) and  
5 incubated overnight at 4°C. Afterward, the plates were washed three times with PBS-  
6 tween 20 (50mM tris,100mM sodium chloride, 0.05% tween 20 and pH 7.4) using an  
7 automatic microplate washer (Immunowash 1575, Biorad, USA) followed by block of  
8 non-specific immune sites with carbonate buffer plus 8% nonfat dry milk and  
9 incubation of 1h at 37°C. Control and test sera samples were diluted 1:20 in PBS-  
10 tween 20 plus 5% nonfat dry milk and 0.1ml of this mixture were added to the  
11 microtiter plates in duplicate and incubated at 37°C for 1 h. After another thrice of  
12 rinsing, 0.1 ml of peroxidase-labeled anti-chicken IgY antibody (A9046, Sigma-  
13 Aldrich, USA), diluted 1:500 in PBS-tween 20 plus 5% nonfat dry milk, was added in  
14 each well and incubated at 37°C for 1 h.

15           Plates were washed thrice and the peroxidase activity was revealed by  
16 adding 0.1 ml of tetramethylbenzidine solution (TMB Single Solution, Invitrogen,  
17 USA). The reaction was stopped by adding 0.05 ml of 1 N HCl and the optical density  
18 (OD) was read at 450nm in an ELISA microplate reader (iMark Microplate  
19 Absorbance Reader, Biorad, USA). The same positive and negative control sera  
20 were included in all plates and a corrected OD value was calculated for each sample  
21 as previously described by Garcia et al. (Garcia et al., 2006) to control plate-to-plate  
22 variation. Serum samples were considered positive when  $OD_{corrected} > [OD \text{ mean}$   
23  $(\text{from negative control sera}) + 3SD$  (standard deviation from negative control sera  
24 from all plates)].

25

## 1 2.4 DNA extraction and PCR

2 For each animal, 0.5 g of brain tissue was homogenized and DNA extraction  
3 was performed using a commercial kit (PureLink Genomic DNA Mini Kit, Invitrogen,  
4 USA) following the manufacturer's instructions. DNA samples were stored at -20°C  
5 until molecular analysis.

6 PCR reactions were performed to detect *N. caninum* DNA in brain tissue  
7 from birds. Nc5 gene was targeted as previously described by Müller et al. (1996)  
8 with minor modification. PCR reaction contained 2µl of genomic DNA, 1x PCR buffer,  
9 0.2mM of each dNTP, 2.5mM MgCl<sub>2</sub>, 1.25 U of Taq polymerase (Platinum Taq DNA  
10 Polymerase, Invitrogen, USA), 0.8µM of each primer (Np21-plus and Np6plus), in a  
11 final solution of 25µl. Reactions was carried out in an automatic DNA thermal cycler  
12 (Veriti Thermal Cycler, Applied Biosystems, USA) following the conditions: 94°C for 5  
13 min for initial denaturation, 35 cycles of 94°C for 30 seg, 63°C for 30 seg, 72°C for 1  
14 min followed by a final extension step at 72°C for 7 min. Positive and negative  
15 controls, constituted of DNA from tachyzoites of Nc-1 strain and ultra-pure water  
16 respectively, were included in all analyses. PCR products were analyzed by  
17 electrophoresis through a 1.5% agarose gel stained with Sybr Safe DNA Gel Stain  
18 (Invitrogen, USA) and visualized under UV light. A 100 bp DNA ladder (Invitrogen,  
19 USA) was used in all agarose gel.

20

## 21 2.5 Statistical analysis

22 All the variables were analyzed by means of the chi-square test or Fisher's  
23 exact test, using the Epi Info software, version 6.04b (DEAN et al., 1994). P-values ≤  
24 0.05 were considered to be significant.

25

### 1 3. Results

2  
3 Out of 249 samples analyzed, 79 (31.7%) were considered to be positive by  
4 ELISA test. Regarding the sex of the doves, there was no significant statistical  
5 difference ( $p>0.05$ ), with 33.3% (38/114) of males positive and 30.3% (41/135) of  
6 females positive at the test. There was a statistically difference between the capture  
7 site ( $p<0.05$ ), since positivity was higher between doves from the soybean seed plant  
8 (38.6%, 54/140), compared with those from the university campus (23.7%, 19/80)  
9 and the dairy cattle farm (20.7%, 6/29) (Table 1). Average optical density from  
10 positive samples was  $0.80\pm 0.26$ , while for negative sample was  $0.18\pm 0.12$ . Optical  
11 density obtained by ELISA test of each animal from different capture site is plotted in  
12 Fig. 1. Serum samples with optical density equal or higher than 0.433 were  
13 considered to be positive.

14 According by nested PCR for PCR for Nc5 gene, no specific DNA fragment  
15 for *N. caninum* was observed in none of brain samples.

### 17 4. Discussion

18  
19 To the best of the author's knowledge, this is the first study that reports an  
20 occurrence of antibodies against *N. caninum* in doves of *Z. auriculata* species, where  
21 a seroprevalence of 31.72% was observed. A previously study with eared doves has  
22 already been done, but different from us, the authors used IFA for *N. caninum*  
23 antibody detection, however, none animal was positive (Mineo et al., 2011).  
24 However, there are some reports showing that birds present an immune response  
25 against *N. caninum* infection (Costa et al., 2008; Gondim et al., 2010; Martins et al.,

1 2011). In all previously studies, serological tests were performed using either IFAT or  
2 MAT, however here, an ELISA test was optimized and used for the first time for  
3 detection of antibodies against *N. caninum* in birds.

4 It was not observed statistical difference between the sex of birds and  
5 seropositivity against *N. caninum*. Similar results were found in studies with *T. gondii*,  
6 where no difference serology considering sex of doves (*Z. auriculata*) and pigeon  
7 (*Columba livia*) was found (Barros et al., 2014; Tsai et al., 2006). Regarding the  
8 capture site, there was a statistically significant difference with higher number of  
9 positive animals captured in a soybean seed plant. Presence of dogs in some areas  
10 could be a reason for this difference, however, doves are able to fly up to 117 km  
11 daily from breed colonies (Bucher and Bocco, 2009), then, they can also be infected  
12 in an area different from the capture site. Although small number of doves from dairy  
13 cattle farm was analyzed, positive results were found, what could contribute to  
14 increase of reproductive problems, since previously studies have shown that  
15 presence of birds on dairy farm leads to higher risk for abortions associated to *N.*  
16 *caninum* (Bartels et al., 1999).

17 Birds are used as a good bioindicator of environmental contamination since  
18 they usually fed on the ground, being exposed to several pathogens, including *N.*  
19 *caninum* oocysts. Studies performed before showed that birds, as chickens, dove,  
20 pigeon among others, are source of infection of *T. gondii* (Barros et al., 2014; Pena  
21 et al., 2013), *Cryptosporidium* spp. (Li et al., 2015) and *Salmonella* spp. (Cano-  
22 Terriza et al., 2015).

23 Costa et al (2008) were pioneers in using serological test to study  
24 immunological response against *N. caninum* in birds. Using chickens as a model and  
25 by means of IFAT with a cut off of 50 they observed an overall seroprevalence of

1 12.5%, however, when they compared the origin of these animals, outdoor chickens  
2 had higher seropositive samples compared with indoor group, evidencing that  
3 animals that have access to different environmental conditions may be more exposed  
4 to this parasite. Afterwards, others studies with chickens were performed, with  
5 seroprevalence ranging from 5.7% to 83.6% (Camillo et al., 2015; Gonçalves et al.,  
6 2012; Martins et al., 2011).

7       Antibodies against *N. caninum* have also been described in other bird species  
8 worldwide (Mineo et al., 2011; Molina-López et al., 2012; Salant et al., 2015).  
9 Seroprevalence studies with the common raven (*Corvus corax*) from Spain showed  
10 that 35.8% were positive (Molina-López et al., 2012) while 16.4% of ravens (*C.*  
11 *cornix*, *C. monedula* and *C. splendens*) from Israel were positive (Salant et al., 2015).  
12 This difference in the serological prevalence may be due to the species of bird  
13 analyzed, geographical region studied and serological and cut off used. In our study,  
14 we used an ELISA test, which is more sensitive than IFAT, and could explain a  
15 higher prevalence in this study than reported by Mineo et al (2011) that used IFAT to  
16 evaluate the presence of antibodies against *N. caninum* in doves *Z. auriculata*.

17       Regarding to molecular diagnostic, no DNA amplification was observed in all  
18 samples tested. It shows that probably this species of doves do not bear a chronic  
19 infection of *N. caninum* with no development of cyst in brain tissues. Du et al. (2015)  
20 reported for the first time that pigeons can be the intermediate host for *N. caninum*  
21 since DNA from the parasite was observed in brain samples, although no species  
22 name from these pigeons was mentioned. Other studies have been reported DNA of  
23 *N. caninum* in brain samples from birds, including chickens (*Gallus domesticus*)  
24 (Costa et al., 2008), sparrow (*Passer domesticus*) (Abdoli et al., 2015; Gondim et al.,  
25 2010), hooded crow (*C. cornix*), western jackdaw (*C. monedula*) (Salant et al., 2015),

1 magpies (*Pica pica*) and common buzzard (*Buteo buteo*) (Darwich et al., 2012).  
2 McGuirre et al. (1999) observed positive PCR results in pigeons (*Columba livia*)  
3 inoculated with different doses of *N. caninum* tachyzoites, even for those who was  
4 seronegative after 6 weeks post inoculation, showing that PCR can detect chronic *N.*  
5 *caninum* infection. Mineo et al. (2011) have been detected *N. caninum* in tissue from  
6 wild and captive birds, since positive immunostaining of cysts tissue with anti-*N.*  
7 *caninum* polyclonal antibodies were observed in the cloacae musculature of a red-  
8 and-green macaw (*Ara chloropterus*) and in the cervical musculature of a blue-  
9 fronted Amazon parrots (*Amazona aestiva*). The same authors have also detected  
10 Apicomplexa-like tissue cysts in the pectoral muscle of a pigeon (*C. livia*) however  
11 immunohistochemistry was not able to detect *N. caninum* antigens.

12

## 13 **5. Conclusions**

14

15 The present study reported for the first time antibodies against *N. caninum* in  
16 doves *Z. auriculata* naturally infected. We conclude that doves have contact with *N.*  
17 *caninum* leading to an immunological response but molecular detection was not  
18 observed, indicating that species of bird may be not a good host for this parasite,  
19 however further studies should be performed in order to evaluate the real importance  
20 of this doves in the epidemiologic cycle of the disease.

21

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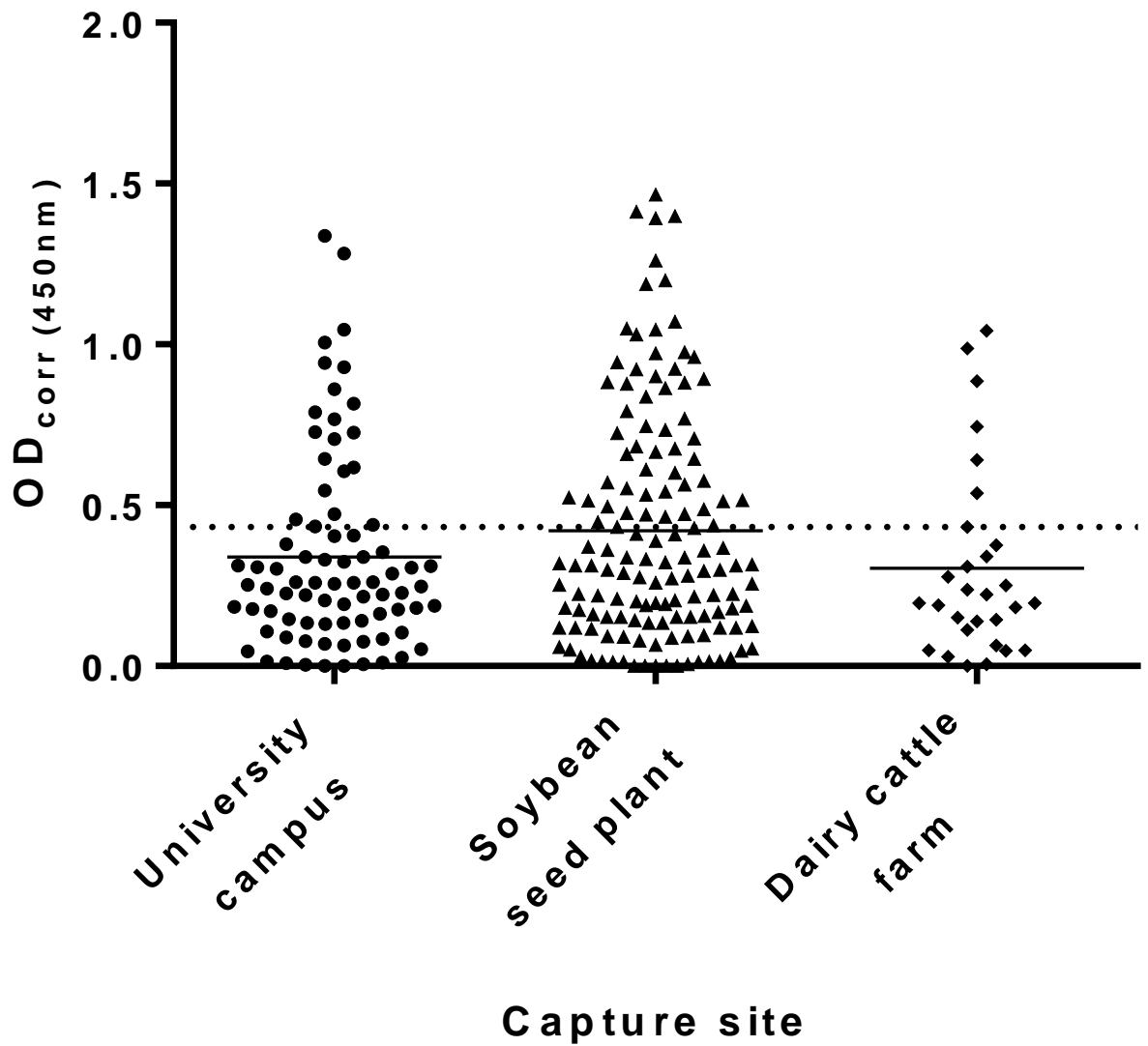
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**Table 1.** Association of sex and capture site with the presence of antibodies against *N. caninum* by an ELISA test in eared doves (*Zenaida auriculata*) from Southern Brazil.

<b>Variables</b>	<b>Positive<sup>a</sup> (%)</b>	<b>Negative (%)</b>	<b>Total</b>	<b>p-value</b>
<b>Sex</b>				
Male	38 (33.3)	76 (66.7)	114 (45.7)	0.71
Female	41 (30.3)	94 (69.7)	135 (54.3)	
Total	79 (31.7)	170 (68.3)	249 (100)	
<b>Capture site</b>				
Soybean seed plant	54 (38.6)	86 (61.4)	140 (56.2)	0.03
University campus	19 (23.7)	61 (76.3)	80 (32.1)	
Dairy cattle farm	6 (20.7)	23 (79.3)	29 (11.7)	
Total	79 (31.7)	170 (68.3)	249 (100)	



**Fig. 1.** Serological result by an ELISA test for *N. caninum* in eared doves (*Zenaida auriculata*) trap-captured from different location in Southern Brazil. Dashed line indicates the positive cut off (0.433) and line the mean.

1 **5 ARTIGO PARA PUBLICAÇÃO II\*\***

2

3 **EXPERIMENTAL INOCULATION OF *NEOSPORA CANINUM* TACHYZOITES IN**  
4 **EARED DOVES (*ZENaida AURICULATA*)**

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## 1 **Abstract**

2 *Neospora caninum* is an apicomplexan parasite with worldwide distribution. Although  
3 a positive association between the presence of birds and abortions in cattle associate  
4 to *N. caninum* has been reported, the role of the birds in the epidemiologic cycle of  
5 the parasite is unknown. Since there are no studies evaluating *N. caninum* in birds of  
6 *Zenaida auriculata* species, the present study aimed to determine if eared doves (*Z.*  
7 *auriculata*) can act as intermediate host for *N. caninum*. Eighteen birds were divided  
8 in four groups, G1, G2 and G3 and received  $2 \times 10^6$  tachyzoites of NC-1 strain by  
9 different routes, being subcutaneous, intramuscular and intraperitoneal respectively.  
10 G4 was composed by three animals and remained as negative control. Serum  
11 samples were collected weekly and one animal from G1, G2 and G3 were euthanized  
12 on the 7<sup>th</sup> and 14<sup>th</sup> day post inoculation (d.p.i.). The remaining birds were euthanized  
13 after 28 d.p.i.. After euthanasia and necropsy, tissues from the doves were submitted  
14 for histopathology, PCR and dog bioassay to detect the parasite. Dogs were fed with  
15 tissue from birds and monitored for 30 days. Serum samples were collected weekly  
16 for serological analysis and feces samples were collected daily until the end of the  
17 experiment for coproparasitological examinations. No dove showed clinical signs of  
18 infection, but all of them seroconverted after inoculation, with stronger immunological  
19 response in birds from G3. Positive PCR was found in lung tissue of one animal from  
20 G3 euthanized on the 7<sup>th</sup> d.p.i. and an inflammatory infiltrate was observed in the  
21 histopathology in the lung and kidney from these dove. No dogs shed oocysts or  
22 seroconverted. Our results indicate that intraperitoneal route induced infection in  
23 doves but the parasite may have been eliminated by the host, being doves resistant  
24 to chronic infection.

25

26 **Keywords:** *Neosporosis*, NC-1 strain, *Zenaida auriculata*, birds, dogs bioassay

27

## 28 **1. Introduction**

29

30 *Neospora caninum* is a protozoan parasite that was first described by Bjerkås  
31 et al (1984) in dogs who presented cysts associated with lesion in the central nervous  
32 system and skeletal muscles. Canids, including domestic dog, Australian dingo  
33 (*Canis familiaris*), coyote (*Canis latrans*) and gray wolf (*Canis lupus*) are definitive  
34 hosts of the parasite (Dubey et al., 2011; Gondim et al., 2004; King et al., 2010;

1 McAllister et al., 1998). Infection by *N. caninum* has been reported in many animals,  
2 including birds, however the zoonotic potential is unknown since just serological  
3 presence of antibodies against the parasite has been described in humans (Darwich  
4 et al., 2012; Dubey and Schares, 2011; Tranas et al., 1999).

5 Neosporosis is an important disease in cattle, being responsible for high  
6 economic losses worldwide due to reproductive problems such as abortion and  
7 persistently infected calves. In Brazil, economic losses are estimated in 153 million  
8 dollars per annum, mainly due to beef industry, which contributes to lower economic  
9 growth of the country (Reichel et al., 2013). Although the disease is important in  
10 dogs, since it can lead to neuromuscular problems, clinical cases are rare and  
11 isolated (Patitucci et al., 1997). In birds, neosporosis is not fully understood, however  
12 previously studies have demonstrated a positive correlation between the presence of  
13 birds and high seroprevalence in cattle, suggesting that birds could play an important  
14 role in the epidemiology of the disease in these animals (Bartels et al., 1999; Otranto  
15 et al., 2003).

16 The dove *Zenaida auriculata* is a bird from Columbiformes order, which  
17 includes other pigeon and dove species. These animals are native from Caribbean to  
18 South America, habiting mainly regions with semiarid climates, widely distributed  
19 throughout Brazilian territory (Souza et al., 2007). In some countries from South  
20 America, such as Argentina, Colombia and Brazil, the population of doves has  
21 increased out of control, being considered as a pest in the agriculture due to damage  
22 to crops (Bucher and Ranvaud, 2006). Although present during all year in some  
23 regions from Brazil, the population density of *Z. auriculata* is higher in the urban area,  
24 which contributes to damage to public environment and transmission of disease to  
25 human beings (Shibatta et al., 2009). Doves are also easily preyed upon by dogs,  
26 which could contribute to the life cycle of the parasite and disseminating of the  
27 disease.

28 The aim of the present study was to evaluate the infection of *N. caninum* in  
29 eared doves (*Z. auriculata*) and the potential of these birds to act as intermediate  
30 hosts of the parasite.

## 33 **2. Material and methods**

34

## 2.1 Animals

Eighteen eared doves (*Z. auriculata*), including 7 males and 11 females, trap captured in Londrina city (23°08'47" to 23°55'46"S/50°52'23" to 51°19'11"W), Paraná state, Southern Brazil, between September 2015 and October 2015 were used to experimental study. All doves were tested to the presence of antibodies against *N. caninum* by ELISA test and all were considered to be negative. Four mongrel dogs, three female and one male, aged 5 months old and healthy were used to bioassay. All dogs were seronegative to *T. gondii*, *Ehrlichia canis*, *N. caninum* and *Leishmania* spp. The dogs were also vaccinated against canine distemper, adenovirus type 1 and type 2, parainfluenza, parvovirus, leptospirosis and rabies (Nobivac<sup>®</sup> Canine, MSD Animal Health, USA) prior to experiment. The animals received anthelmintic treatment with praziquantel, febantel, pyrantel pamoate and ivermectin (Canex Premium, Ceva Santé Animale, Brazil), were fed with commercial feed, received distilled water *ad libitum* and were allocated separately in individual stalls.

All procedures involving animals were approved by Animal Ethics Committee of State University of Londrina (CEUA no. 157/15) and the Brazilian Institute of the Environment (IBAMA-SISBIO no. 16.428-1)

## 2.2 *Neospora caninum* strain

*Neospora caninum* 1 (NC-1) strain (Dubey et al., 1988) were used to produce the inoculum and for preparation of antigens for serological test. Tachyzoites were grown in Vero cells supplemented with RPMI medium, bovine fetal serum and penicillin (10,000 U) and streptomycin (10mg/ml). After five days of inoculation, most of tachyzoites were free in the flask and cells were scraped and passed through 27G needles. Then, the solution was centrifuged 2,000xg for 10 min and the supernatant was discarded. The resultant pellet was resuspended in 1 ml of RPMI solution and tachyzoites were counted using a Neubauer chamber.

## 2.3 Experimental design

The doves were divided into 4 groups, group 1 (G1, n=5), group 2 (G2, n=5), group 3 (G3, n=5) and group 4 (G4, n=3). The G1, G2 and G3 received the same

1 inoculum concentration ( $2 \times 10^6$  tachyzoites) by different routes, being subcutaneous,  
2 intramuscular and intraperitoneal respectively. The G4 was the negative control and  
3 received PBS solution by intraperitoneal route.

#### 4 5 2.4 Sample collection

6  
7 Blood samples from the birds were collected by brachial venipuncture at days  
8 0, 7, 14, 21 and 28 and serum samples obtained were stored at  $-20^{\circ}\text{C}$  until  
9 serological analysis. On day 7<sup>th</sup> and 14<sup>th</sup>, one animal of G1, G2 and G3 were  
10 euthanized and tissue samples (brain, heart, liver, lung, kidney and pectoral muscle)  
11 were collected for histopathological and PCR analysis. All other animals were  
12 euthanized on day 28<sup>th</sup> and also had their tissue samples collected for histopathology  
13 and molecular analysis. All animals were euthanized following the guidelines  
14 established by the Nacional Council for Animal Experimental Control  
15 (CONCEA/Brazil).

#### 16 17 2.5 Enzyme-linked immunosorbent assay test (ELISA)

18  
19 In order to detect antibodies IgG against *N. caninum*, an enzyme-linked  
20 immunosorbent assay test (ELISA) was used. For the preparation of antigen,  
21 tachyzoites of NC-1 strain were growth in Vero cell supplemented with RPMI  
22 medium, bovine fetal serum and penicillin (10,000 U) and streptomycin (10mg/ml).  
23 Subsequent, the cells were scraped and passed through 27G needles. The resultant  
24 solution was centrifuged 10,000 g for 10 min and pellet was mixed with ceramic  
25 beads through vortex three times of 10 minutes each. The solution was centrifuged  
26 again following the same conditions described before and the pellet was discarded.  
27 Antigen solution was submitted to protein concentration measure by using a  
28 commercial kit (Pierce™ BCA, Thermo Fisher, USA). The antigen was stored at  $-20^{\circ}$   
29 C until performance of serological tests.

30 Previously tests were conducted to determine the optimal antigen  
31 concentration and dilutions of serum and conjugate. Flat-bottom 96 well polystyrene  
32 microtitration plates (Nunc-Immuno Plate, MaxiSorp, Denmark) were coated with  
33 0.1ml of crude antigen (2.5 $\mu\text{g}/\text{ml}$ ) diluted in 0.1M carbonate buffer (pH 9.6) and  
34 incubated overnight at  $4^{\circ}\text{C}$ . Afterward, the plates were washed three times with PBS-

1 tween 20 (50mM tris,100mM sodium chloride, 0.05% tween 20 and pH 7.4) using an  
2 automatic microplate washer (Immunowash 1575, Biorad, USA) followed by block of  
3 non-specific immune sites with carbonate buffer plus 8% nonfat dry milk and  
4 incubation of 1h at 37°C. Control and test sera samples were diluted 1:20 in PBS-  
5 tween 20 plus 5% nonfat dry milk and 0.1ml of this mixture were added to the  
6 microtiter plates in duplicate and incubated at 37°C for 1 h. After another thrice of  
7 rinsing, 0.1 ml of peroxidase-labeled anti-chicken IgY antibody (A9046, Sigma-  
8 Aldrich, USA), diluted 1:500 in PBS-tween 20 plus 5% nonfat dry milk, was added in  
9 each well and incubated at 37°C for 1 h.

10 Plates were washed thrice again and the peroxidase activity was revealed by  
11 adding 0.1 ml of tetramethylbenzidine solution (TMB Single Solution, Invitrogen,  
12 USA). The reaction was stopped by adding 0.05 ml of 1 N HCl and the optical density  
13 (OD) was read at 450nm in an ELISA microplate reader (iMark Microplate  
14 Absorbance Reader, Biorad, USA). The same positive and negative control sera  
15 were included in all plates and a corrected OD value was calculated for each sample  
16 as previously described by Garcia et al. (2006) to control plate-to-plate variation.  
17 Serum samples were considered positive when  $OD_{corrected} > [OD \text{ mean (from}$   
18  $\text{negative control sera) + 3SD (standard deviation from negative control sera from all}$   
19  $\text{plates)]}$ .

20

## 21 2.6 DNA extraction and PCR

22

23 A total of 0.25 mg of each tissue sample from each animal was submitted to  
24 DNA extraction using a commercial kit (PureLink Genomic DNA Mini Kit, Invitrogen,  
25 USA) following the manufacturer's instructions. DNA samples were stored at -20° C  
26 until molecular analysis.

27 PCR reactions were performed to detect *N. caninum* DNA in all tissue  
28 samples from birds collected during the necropsy. Nc5 gene was amplified as  
29 previously described by Müller et al. (1996) with minor modification, which amplifies a  
30 fragment of 337 bp. PCR reaction contained 2µl of genomic DNA, 1x PCR buffer,  
31 0.2mM of each dNTP, 2.5mM MgCl<sub>2</sub>, 1.25 U of Taq polymerase (Platinum Taq DNA  
32 Polymerase, Invitrogen, USA), 0.8µM of each primer (Np21-plus and Np6plus), in a  
33 final solution of 25µl. Reactions was carried out in an automatic DNA thermal cycler  
34 (Veriti Thermal Cycler, Applied Biosystems, USA) following the conditions: 94°C for 5

1 min for initial denaturation, 35 cycles of 94°C for 30 seg, 63°C for 30 seg, 72°C for 1  
2 min followed by a final extension step at 72°C for 7 min. Positive and negative  
3 controls, constituted of DNA from tachyzoites of Nc-1 strain and ultra-pure water  
4 respectively, were included in all analyses. PCR products were analyzed by  
5 electrophoresis through a 1.5% agarose gel stained with Sybr Safe DNA Gel Stain  
6 (Invitrogen, USA) and visualized under UV light. A 100 bp DNA ladder (Invitrogen,  
7 USA) was used in all agarose gel.

## 8 9 2.7 Histopathology

10  
11 All tissue samples collected during the necropsy of birds were kept in a 10%  
12 buffered formalin solution for 24 h, transferred subsequently to a 70% ethyl alcohol  
13 solution and embedded in paraffin. Paraffin-embedded tissues were serially  
14 sectioned at 4 µm thick and stained with hematoxylin-eosin (HE) for histopathological  
15 evaluation.

## 16 17 2.8 Bioassay in dogs

18  
19 The dogs (n=4) were fed with a pool of tissue of approximately 70g (liver, lung,  
20 brain, heart, kidney and pectoral muscle) from inoculated birds. Each dog received a  
21 pool of all birds from different groups (i.e., dog 1 received tissue from all birds from  
22 G1, dog 2 birds from G2, dog 3 from G3 and dog 4 from G4). Physical examination of  
23 the dogs was performed every day during the experimental period. Blood samples  
24 from the dogs were collected by cephalic venipuncture at day 0 and every 7 days  
25 until the end of the experiment. Fecal samples were collected daily until 30<sup>th</sup> day after  
26 dogs were fed with tissue for coproparasitological examinations. During the  
27 experimental period, stalls were cleaned and disinfected every day with sodium  
28 hypochlorite at 2%.

## 29 30 2.9 Indirect fluorescent antibody test (IFAT)

31  
32 Serum samples of all dogs were tested for antibodies IgG against *N. caninum*  
33 by an indirect fluorescent antibody test (IFAT) according with technique previously  
34 described (Dubey et al., 1988). Anti-dog IgG (F4012, Sigma-Aldrich, USA)

1 conjugated with fluorescein isothiocyanate were used as secondary antibodies.  
2 Samples with titer  $\geq 50$  were considered positive.

### 3 4 2.10 Coproparasitological examinations

5  
6 Fecal samples from dogs were examined daily during a period of 30 days by  
7 sucrose flotation technique (density of  $1.2 \text{ g/cm}^3$ ) for detection of *Neospora*-like  
8 oocysts as previously described with minor modification (Sheather, 1923). The total  
9 volume of feces from each day and each animal were homogenized and weighed.  
10 Five grams from this homogenized sample were diluted in distilled water, filtered with  
11 gauze and centrifuged at 1,200 g for 10 min. The pellet was mixed with sucrose  
12 solution in a 15ml plastic tube and centrifuged at 1,200 g for 10 minutes. Afterwards,  
13 25  $\mu\text{l}$  from the upper meniscus of the solution was recovered and transferred to a  
14 slide and covered with a coverslip for microscopic examination.

## 15 16 17 **3. Results**

18  
19 All inoculated birds did not showed any clinical signs or died after infection.  
20 The doves from all groups seroconverted after challenge, with a peak on 7 d.p.i.,  
21 however OD decrease during the experiment period, being no detectable in one  
22 animal inoculated through by subcutaneous and two by intramuscular route, while all  
23 doves inoculated by intraperitoneal route were considered to be positive (Fig. 1). All  
24 birds from control group remained negative through experiment. The OD mean of G3  
25 ( $1.589 \pm 0.79$ ) were higher than G1 ( $0.779 \pm 0.71$ ) and G2 ( $0.723 \pm 0.52$ ).

26 One dove from G3, euthanized 7 d.p.i., had lung tissue positive by means of  
27 PCR. All of other samples were negative by molecular tests. In the histopathology  
28 evaluation, an inflammatory focus of mononuclear leukocytes was observed in the  
29 lung and kidney of the same bird that was positive in PCR, however no cyst  
30 compatible with *N. caninum* was observed in all samples analyzed.

31 In the bioassay in dogs, no oocysts were detected in the dog feces during the  
32 examination. Furthermore, none of the dogs showed any signs of clinical infection or  
33 seroconversion after fed with tissue from inoculated birds. All results are summarized  
34 in table 1.

#### 1 4. Discussion

2  
3 In the present study no eared dove showed signal clinical of infection or  
4 mortality, being considered as resistant to neosporosis. Previously studies have  
5 shown that other species from birds, such as quails (*Coturnix coturnix japonica*) (de  
6 Oliveira et al., 2013) and chickens (*Gallus gallus domesticus*) (Furuta et al., 2007;  
7 McGuire et al., 1999; Munhoz et al., 2014) are resistant to *N. caninum* infection by  
8 NC-1 tachyzoites, however Mineo et al. (2009) reported that pigeon from *Columba*  
9 *livia* species are susceptible, with dissemination of the parasite through lungs, heart,  
10 liver, spleen, kidney and central nervous system. This difference in susceptibility to *N.*  
11 *caninum* infection could be related to different biological models and difference in  
12 inoculum concentration, since they used a higher number ( $10^7$ ) of parasites.

13 All doves had antibodies above the cut off seven days after inoculation,  
14 however, the OD mean from all group decreases during the experimental period. This  
15 is in agreement with studies performed by de Oliveira et al. (2013) and Mineo et al.  
16 (2009), who showed that birds inoculated with tachyzoites present a peak of  
17 antibodies usually 14 days post inoculation but rapidly decreases, being undetectable  
18 after 30 days by means of IFAT. Animals from G3, that received tachyzoites by  
19 intraperitoneal route, had higher OD in ELISA test. It corroborates previously studies  
20 that reported intraperitoneal inoculation of *N. caninum* tachyzoites induce stronger  
21 immune response compared mainly with subcutaneous inoculation (de Oliveira et al.,  
22 2013; McGuire et al., 1999; Mineo et al., 2009).

23 According with PCR analysis, only lung sample from one bird of G3 was  
24 positive. It seems that intraperitoneal inoculation is a better route to parasite  
25 dissemination in eared doves, what have been reported in pigeons (*C. livia*), where  
26 birds inoculated by intraperitoneal route had positive results in different techniques,  
27 as tissue culture, PCR and histopathology (McGuire et al., 1999). In our study, the  
28 dove that was positive in PCR was euthanized 7 days post inoculation, which shows  
29 that *N. caninum* disseminate during the first days after infection, being undetectable  
30 after 4 weeks, showing that eared doves are resistant to chronic infection with *N.*  
31 *caninum* tachyzoites, as reported in quails and chickens (de Oliveira et al., 2013;  
32 Furuta et al., 2007).

33 Only one dove showed alterations in the histopathology evaluation that could  
34 be associated with neosporosis, with presence of mononuclear inflammatory focus. A

1 mononuclear inflammatory infiltrate was also observed in livers from quails 3 and 7  
2 d.p.i. with *N. caninum* tachyzoites. McGuire et al. (1999) observed typical lesions of  
3 neosporosis in one pigeon (*C. livia*) inoculated with higher dose of tachyzoites,  
4 although they used NC-2 and NC-Liverpool, different strains from the our study, what  
5 could explain this difference in the pathogenesis of chronic infection, since these  
6 alterations were observed in animals euthanized six weeks post inoculation. No  
7 lesions were observed in doves euthanized after 28 d.p.i. and similar results were  
8 observed by de Oliveira et al. (2013) and Munhoz et al. (2014), who did not find  
9 changes in histopathology of chronically infected birds.

10 In the bioassay, during all experimental period, no dogs showed clinical signs  
11 of infection, shed oocysts or seroconverted after fed with tissue from doves  
12 inoculated with *N. caninum*. Other studies have also reported that dogs fed with  
13 tissue from birds inoculated with NC1- strain did not shed oocysts or seroconverted  
14 (de Oliveira et al., 2013; Munhoz et al., 2014). These results suggest that *N. caninum*  
15 cysts were no present in tissues from doves and probably infection with NC-1  
16 tachyzoites, different from other species, is not a good model to induce chronic  
17 infections in birds, as reported in quails and chickens (de Oliveira et al., 2013;  
18 Gondim et al., 2002; Munhoz et al., 2014).

## 20 5. Conclusions

21  
22 Based on results presented in this study, we conclude that intraperitoneal  
23 route induced infection in doves (*Z. auriculata*), but these animals are not a good  
24 intermediate host for *N. caninum*, since it seems like the parasite may have been  
25 eliminated by the host, being considered resistant to chronic infection. Further  
26 studies, using oocysts as infection model are necessary to evaluate the course of  
27 infection and determine the importance of doves in the epidemiological cycle of  
28 neosporosis.

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**Table 1.** Detection of *Neospora caninum* by different test in eared doves (*Zenaida auriculata*) inoculated with tachyzoites of NC-1 strain.

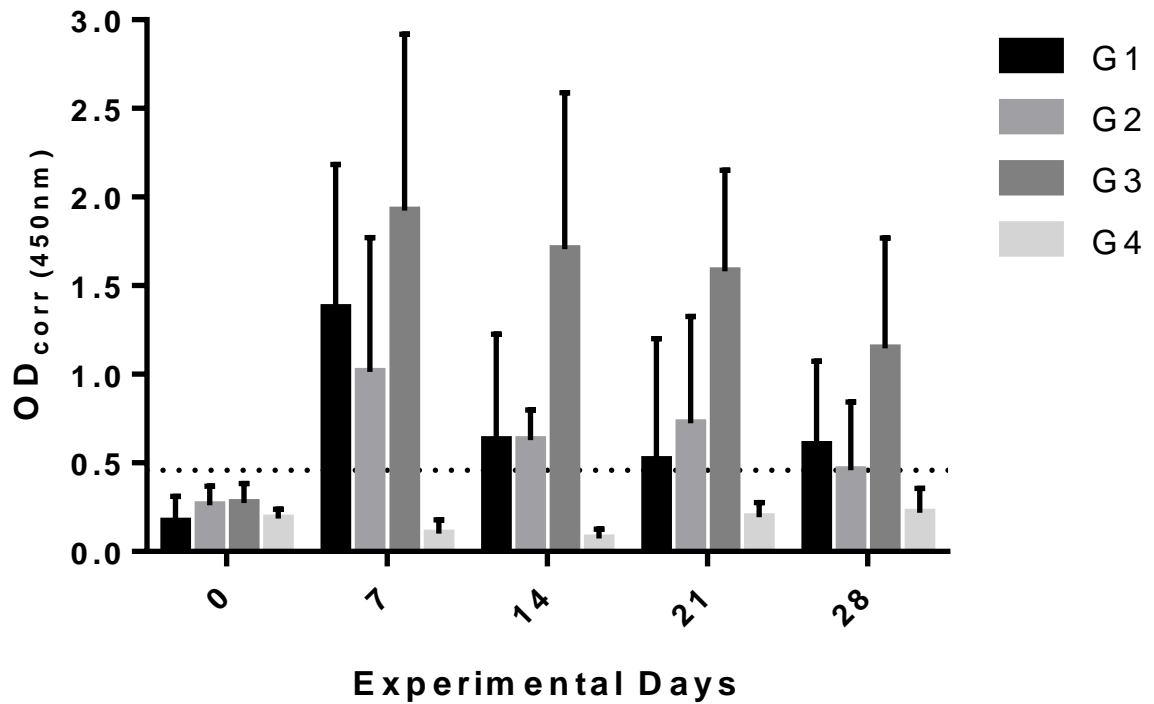
Group	Dose	Via <sup>a</sup>	OD mean <sup>b</sup>	PCR	HP <sup>c</sup>	Bioassay in dogs	
						Oocyst shedding	IFAT titer
G1	2x10 <sup>6</sup>	SC	0.779	0/5	-	-	0
G2	2x10 <sup>6</sup>	IM	0.723	0/5	-	-	0
G3	2x10 <sup>6</sup>	IP	1.589	1/5	+	-	0
G4	na	IP	0.146	0/5	-	-	0

n.a.: not applicable.

a: SC: subcutaneous; IM: intramuscular; IP: intraperitoneal

b: Optical density obtained by ELISA test

c: Histopathological examination



**Fig 1.** Immunological response of eared doves (*Zenaida auriculata*) inoculated with *Neospora caninum* tachyzoites evaluated by the indirect enzyme-linked immunosorbent assay (ELISA). All birds received the same parasite concentration by different routes, being G1 subcutaneous, G2 intramuscular and G3 intraperitoneal route (bars= OD mean from animals and error bars= standard deviation). Dashed line indicates positive cut off (OD=0.4579)

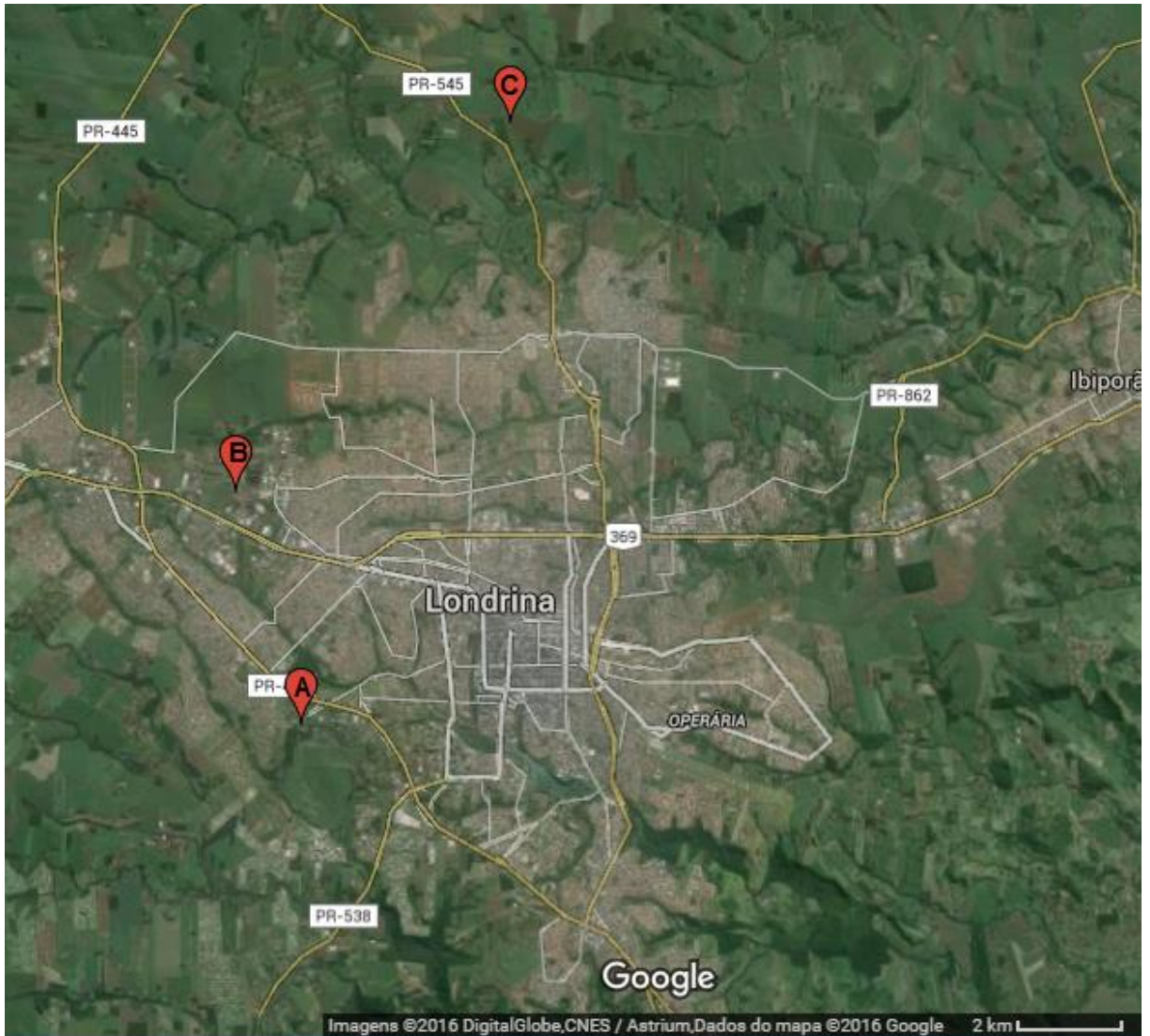
## 6 CONCLUSÃO

- Foi observado uma ocorrência de anticorpos contra *N. caninum* de 31.72%, em pombos da espécie *Z. auriculata*, sendo o primeiro relato da presença de anticorpos contra *N. caninum* nessa espécie de pombo.
- Pombos inoculados com taquizoítos da cepa NC-1 de *N. caninum* apresentaram uma elevada resposta imunológica 7 dias após inoculação
- Inoculação intraperitoneal de taquizoítos da cepa NC-1 de *N. caninum* foi capaz de induzir infecção em pombos, embora eles foram resistentes à infecção crônica.
- Cães alimentados com tecidos de pombos experimentalmente infectados com taquizoítos da cepa NC-1 não eliminaram oocistos ou soroconverteram, mostrando que os pombos da espécie *Z. auriculata* não são hospedeiros intermediários do *N. caninum* nas condições do presente estudo.

**ANEXOS**

## ANEXO A

Mapa do local de captura dos pombos. A: Campus universitário; B: Indústria de processamento de grãos; C: Fazenda de bovinocultura leiteira.



**ANEXO B****Ofício do Comitê de Ética em Experimentação Animal nº 70/2008**

UNIVERSIDADE  
ESTADUAL DE LONDRINA



GOVERNO DO  
PARANÁ

**COMITÊ DE ÉTICA EM EXPERIMENTAÇÃO ANIMAL**

OF. CIRC. CEEA Nº 70/2008

Londrina, 10 de setembro de 2008.

Prezado Pesquisador

O CEEA/UDEL, reunido aos 09 de setembro do ano corrente, avaliou o projeto de pesquisa intitulado "**Caracterização genética de isolados de *Toxoplasma gondii* de pombas (*Zenaida auriculata*) do município de Londrina, Paraná**", registrado no CEEA sob o nº 35/08, desenvolvido sob sua responsabilidade, julgando-o *aprovado* para execução por entender que os princípios éticos postulados pelo Colégio Brasileiro de Experimentação Animal estão respeitados.

Serão utilizados 384 pombos capturados em praças públicas do município de Londrina e 768 camundongos adultos procedentes do Biotério central da UEL.

Tendo em vista o grande número de pombos capturados e o ineditismo e a complexidade desse procedimento, o CEEA/UDEL sugere que outras investigações relevantes sejam realizadas aproveitando material biológico colhido desses animais.

Cumprir orientar que caso se pretendam quaisquer alterações no protocolo experimental aprovado, deve-se submeter o novo protocolo à apreciação do CEEA/UDEL anteriormente à execução das modificações.

Sem mais para o momento, subscrevo-me.

Cordialmente,

A handwritten signature in black ink, appearing to read 'J. A. Naylor Lisboa'.

Prof. Dr. Julio Augusto Naylor Lisboa  
Coordenador do CEEA/UDEL

**Ilmo. Sr.  
Prof. Dr. João Luis Garcia  
Coordenador do Projeto  
Departamento de Medicina Veterinária Preventiva  
Centro de Ciências Agrárias**

## ANEXO C

Ofício da Comissão de Ética no Uso de Animais nº 157/2015



Universidade  
Estadual de Londrina

## COMISSÃO DE ÉTICA NO USO DE ANIMAIS

OF. CIRC. CEUA Nº 157/2015

Londrina, 04 de Agosto de 2015

Prezado Pesquisador,

A CEUA/UEL reunida em 07 de Julho de 2015 avaliou o projeto de pesquisa intitulado "Avaliação da importância de pombos da espécie *Zenaida auriculata* na epidemiologia do *neospora caninum*", registrado sob o processo CEUA nº8676.2015.81, pesquisa do Centro de Ciências Agrárias, desenvolvido sob sua responsabilidade. Esclarecidos os aspectos metodológicos solicitados, o projeto está **aprovado** para execução entendendo-se que os princípios éticos postulados pelo Conselho Nacional de Controle de Experimentação Animal estão respeitados.

Serão utilizados 4 cães sem raça definida e com idade superior à 3 meses; e 24 aves *Zenaida auriculata*, sendo os cães animais abandonados ou doados e as aves capturadas na cidade e área rural por meio de armadilhas tipo arapuca. O projeto tem como objetivo avaliar a importância dos pombos (*Z. auriculata*) como hospedeiro intermediário do *Neospora caninum*. Para tanto, as aves serão divididas em quatro grupos experimentais, sendo um controle (que receberá uma dose de solução fisiológica i.m.) e os outros 3 desafiados com  $1 \times 10^7$  taquizoítos de *N. caninum* pela via subcutânea, intraperitoneal e intramuscular. Quarenta e cinco dias após o desafio, os animais serão eutanasiados por deslocamento cervical e os tecidos avaliados para a presença do parasita por meio da PCR, imunohistoquímica e bioensaio em cães. Os tecidos e demais materiais biológicos dos pombos serão fornecidos aos cães para alimentação para verificar a infectividade dos cistos teciduais e consequente eliminação dos oocistos do parasita por exames coproparasitológicos. As fezes serão coletadas diariamente durante 30 dias após a ingestão dos tecidos dos pombos e também será realizada coleta de sangue semanal. Os protocolos experimentais estão aprovados com previsão para execução em 14 meses.

Cumprе orientar que caso pretendam-se quaisquer alterações no protocolo experimental aprovado, deve-se submeter o novo protocolo à apreciação da CEUA/UEL anteriormente à execução das modificações.

Coloco-me à disposição para quaisquer esclarecimentos que se fizerem necessária. Sem mais para o momento, subscrevo, cordialmente,

Prof. Dr. Waldiceu Aparecido Verri Junior  
Coordenador da CEUA/UEL

Ilmo. Sr.

Prof. Dr. João Luis Garcia

Coordenador do Projeto

Departamento de Medicina Veterinária Preventiva

Centro de Ciências Agrárias

Com cópia para André Junior da Conceição (Chefe da DP-IC/PROPPG), Chefe do Departamento de Medicina Veterinária Preventiva e Diretor(a) do Centro de Ciências Agrárias.

## ANEXO D

Autorização do Instituto Brasileiro do Meio Ambiente e dos Recursos Naturais  
Renováveis – IBAMA nº 16428-1



Ministério do Meio Ambiente - MMA

Instituto Brasileiro do Meio Ambiente e dos Recursos Naturais Renováveis - IBAMA

Instituto Chico Mendes de Conservação da Biodiversidade - ICMBio

Sistema de Autorização e Informação em Biodiversidade - SISBIO

**Autorização para atividades com finalidade científica**

<b>Número:</b> 16428-1	<b>Data da Emissão:</b> 14/07/2008 10:12	<b>Data de Validade:</b> 14/07/2009
<b>Dados do titular</b>		
Registro no Ibama: 2509292	Nome: João Luís garcia	CPF: 605.644.319-15
Título do Projeto: Caracterização genética de isolados de <i>Toxoplasma gondii</i> de Pombas ( <i>Zenaida auriculata</i> ) do município de Londrina, Paraná.		
Nome da Instituição : Universidade Estadual de Londrina		CNPJ: 78.640.489/0001-53

**Observações, ressalvas e condicionantes**

1	As atividades de campo exercidas por pessoa natural ou jurídica estrangeira, em todo o território nacional, que impliquem o deslocamento de recursos humanos e materiais, tendo por objeto coletar dados, materiais, espécimes biológicos e minerais, peças integrantes da cultura nativa e cultura popular, presente e passa da, obtidos por meio de recursos e técnicas que se destinem ao estudo, à difusão ou à pesquisa, estão sujeitas a autorização do Ministério de Ciência e Tecnologia.
2	Esta autorização não exime o titular e a sua equipe da necessidade de obter as anuências previstas em outros instrumentos legais, bem como do consentimento do responsável pela área, pública ou privada, onde será realizada a atividade.
3	Esta autorização não poderá ser utilizada para fins comerciais, industriais, esportivos ou para realização de atividades inerentes ao processo de licenciamento ambiental de empreendimentos. O material biológico coletado deverá ser utilizado para atividades científicas ou didáticas no âmbito do ensino superior.
4	A autorização para envio ao exterior de material biológico não consignado deverá ser requerida por meio do endereço eletrônico <a href="http://www.ibama.gov.br/cities">www.ibama.gov.br/cities</a> . Em caso de material consignado, consulte <a href="http://www.ibama.gov.br/sisbio">www.ibama.gov.br/sisbio</a> - menu Exportação.
5	O titular de licença ou autorização e os membros da sua equipe deverão optar por métodos de coleta e instrumentos de captura direcionados, sempre que possível, ao grupo taxonômico de interesse, evitando a morte ou dano significativo a outros grupos; e empregar esforço de coleta ou captura que não comprometa a viabilidade de populações do grupo taxonômico de interesse em condição in situ.
6	Este documento não dispensa o cumprimento da legislação que dispõe sobre acesso a componente do patrimônio genético existente no território nacional, na plataforma continental e na zona econômica exclusiva, ou ao conhecimento tradicional associado ao patrimônio genético, para fins de pesquisa científica, bioprospecção e desenvolvimento tecnológico.
7	Em caso de pesquisa em Unidade de Conservação Federal, o pesquisador titular deverá contactar a administração dessa unidade a fim de CONFIRMAR AS DATAS das expedições, as condições para realização das coletas e de uso da infra-estrutura da unidade.

**Locais onde as atividades de campo serão executadas**

#	Município	UF	Descrição do local	Tipo
1	LONDRINA	PR	Universidade Estadual de Londrina	Fora de UC

**Atividades X Táxons**

#	Atividade	Táxons
1	Captura de animais silvestres in situ	Zenaida auriculata
2	Coleta/transporte de amostras biológicas in situ	Zenaida auriculata
3	Coleta/transporte de espécimes da fauna silvestre in situ	Zenaida auriculata (*Qtde: 384)

\* Qtde. de indivíduos por espécie/localidade/unidade de conservação, a serem coletados durante um ano.

**Material e métodos**

1	Amostras biológicas (Aves)	Fezes, Ectoparasita, Sangue, Fragmento de tecido/órgão
2	Método de captura/coleta (Aves)	Outros métodos de captura/coleta(Arapucas-Armadas Isca )

**Destino do material biológico coletado**

#	Nome local destino	Tipo Destino
1	Universidade Estadual de Londrina	Pesquisa

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